# HISTOPATHOLOGICAL AND MOLECULAR STUDIES ON CASEOUS LYMPHADENITIS IN SHEEP AND GOATS IN DUHOK CITY, IRAQ

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ABSTRACT: Caseous lymphadenitis is an infectious bacterial disease of sheep and goats caused by *Corynebacterium pseudotuberculosis*. This study is aimed to estimate the frequency of caseous lymphadenitis in sheep and goats and to compare with the results of bacteriological analysis, histopathology and molecular diagnosis of suspected cases. In this study, 22 (2.1%) suspected cases of caseous lymphadenitis were diagnosed clinically from 1090 sheep and goats slaughtered at Duhok abattoir. The study showed grossly affected mediastinal lymph node was congested and highly oedematous compared to other lymph nodes. Histopathological examination showed presence of abscess in the centre of affected mediastinal lymph node. Based on standard bacteriological methods, *Corynebacterium pseudotuberculosis* was the only causative agent isolated from all suspected cases. The confirmation was performed by amplification of the target genes of *Corynebacterium pseudotuberculosis* including *16S rRNA*, *rpoB* and *pld*. The results showed that the *Corynebacterium pseudotuberculosis* is still the main causative agent of CLA in sheep and goats. The results also revealed molecular and histopathological studies together could be used to identify *Corynebacterium pseudotuberculosis* directly from cases of CLA from sheep and goats. Further studies including study of virulence genes and antimicrobial susceptibility together with phylogenetic analyses are recommended to be done in the future for better understanding of the epidemiology of this pathogen in Duhok city.

Key words: CLA, C. pseudotuberculosis, Histopathology, Molecular diagnosis, Sheep and goat, Duhok.

## **INTRODUCTION**

Caseous lymphadenitis (CLA) is an infectious disease that affects small ruminants including sheep and goats and the disease is caused by *Corynebacterium pseudotuberculosis* (*C. pseudotuberculosis*) (Guerrero *et al.* 2018, Parin *et al.* 2018). *C. pseudotuberculosis* is a Gram-positive, pleomorphic, facultative anaerobic, nonmotile, non-spore forming, non-capsulated, and intracellular microorganism (Guimarães *et al.* 2011). The microorganism is classified into two biovars based on their ability to reduce nitrate and two biovars have been confirmed by biotyping and molecular techniques (Connor *et al.* 2000, Connor *et al.* 2007). The biovar Ovis is unable to reduce nitrate and it mainly affects sheep and goats causing caseous lymphadenitis. The biovar Equi on the other hand, is able to reduce nitrate and is mainly isolated from horses and cattle with ulcerative lymphangitis (Oliveira *et al.* 2016). CLA is characterized by formation of caseous abscesses in lymph nodes and visceral organs as well and it may also cause disorders of reproductive system including orchitis, abortion, and stillbirth (Umer *et al.* 2017). Infection with *C. pseudotuberculosis* is associated with necrosis, inflammation, and enlargement of lymph nodes. In addition, hair loss and chronic abscessation of affected lymph nodes may occur. Sometimes rupture of the affected node followed by pus discharge also is observed (Baird and Fontaine 2007). CLA is one of the economically important diseases in sheep and goats worldwide and the economic losses are due to poor

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growth of wool, decrease in meat, and milk production, reproductive disorders of affected animals and carcasses as well as skin condemnation in abattoirs (Arsenault et al. 2003, Guimarães et al. 2011, Umer et al. 2017). Diagnosis of CLA is based mainly on a post-mortem examination of the lesions, isolation and identification of C. pseudotuberculosis from the pus using phenotypic characterization and biochemical tests (Dorella et al. 2006). The characteristics of the species within the Corynebacterium genus are variable; accordingly advance technique such as polymerase chain reaction (PCR) is used to confirm the detection of the bacterium with high specificity, sensitivity and efficiency. For example, PCR based detection of 16S rRNA (Cetinkaya et al. 2002), RNA polymerase β-subunit (rpoB) (Khamis et al. 2004) and phospholipase D (pld) (Pacheco et al. 2007) genes has been developed to identify C. pseudotuberculosis isolates. The aim of the study was to estimate the frequency of CLA in sheep and goat in Duhok city and to compare with the results of bacterial cultures, histopathology and molecular diagnosis of suspected cases.

# MATERIALS AND METHODS Sample collection

From September to January 2019-2020, 1090 sheep and goat carcasses in a Duhok abattoir were randomly selected for inspection at slaughter house. The samples were taken from the carcasses that had prominent enlargement of the following lymph nodes: submandibular, prescapular, prefemoral, or medistinal. Enlarged abnormal lymph nodes were collected with a sterile scalpel under aseptic conditions and placed in individual sterile containers. The samples were immediately transported in an ice box to the Duhok Research Centre at College of Veterinary Medicine and were processed for microbial isolation, histopathological processing and for molecular study to identify causitve agent(s).

## Histopathology study

Suspected lymph nodes were inspected macroscopically for size, colour, consistency of exudate and presence of lamellate or onion ring morphology. Samples for histopathological study were fixed in 10% neutral buffered formalin (NBF). After 24-48 h the formalin solution was changed then tissue processing was performed manually and sectioning of the tissues was done using a sliding microtome to get 4-5 micrometers thick sections (Leica, Germany). Cut sections were stained with hematoxylin and eosin (H&E) (Luna1968).

# Isolation and identification of bacteria using conventional method

C. pseudotuberculosis is the causative agent of CLA in sheep and goats, for the isolation of the bacterium, pus was collected aseptically to prevent contamination from either surrounding environment or surface of lymph node. Briefly, the surface of abscessed lymph nodes was cleaned with 70% ethanol and incised with a sterile disposable surgical blade. Cheesy substance or pus were collected using sterile disposable swabs and cultured onto 5% blood agar (brain heart infusion agar containing 5% sheep blood). The inoculated plates were incubated for 24-48 h at 37 °C. The plates were checked after 24 h for growth. Standard microbiology methods were used for primary identification and these methods included cultural characteristics, haemolytic activity after 24 and 48h incubation. Colonies with typical morphology of C. pseudotuberculosis were selected and streaked onto blood agar plate. Those isolates were further confirmed by using Gram staining and biochemical tests including catalase, urease and CAMP (Christie, Atkins, Munch-Peterson) tests with Staphylococcus aureus. Suspected C. pseudotuberculosis isolates were then maintained in 50% (v/v) glycerol in brain heart infusion broth (BHIB) (Dorella et al. 2006) and stored till further analysis at -20 °C.

# Molecular detection of *C. pseudotuberculosis* by PCR

## DNA extraction from suspected isolates

PCR was used to confirm the detection of suspected *C. pseudotuberculosis* isolates. To extract DNA, suspected isolates were culturedon 5% sheep blood agar from glycerol stocks. The plates were incubated for 24 h at 37 °C. Thermal lysis method as described by de Sá *et al.* (2013) was used to extract DNA. Briefly 5 to 10 colonies were resuspended in 500  $\mu$ l of TAE buffer and mixed very well. Bacterial suspensions were kept in heating block for 15 min. The suspension was cooled at 4 °C for 5 min then centrifuged at 14,000 x *g* for 2 min. The supernatant was then used as the DNA template for PCR. NanoDrop 2000C spectrophotometer (Thermo Scientific, UK) was used to check DNA was stored at -20 °C.

#### **Primers and PCR conditions**

Primers targeting the 16S rRNA, rpoB and pld genes of C. pseudotuberculosis were considered from previously published work (Çetinkaya et al. 2002, Khamis et al. 2004, Pacheco et al. 2007). Oligonucleotide primers

Gene	Primer name	Sequence (5'—3´)	Product size (bp)	Ref.
16S rRNA	16S-F 6S-R	ACCGCACTTTAGTGTGTGTG TCTCTACGCCGATCTTGTAT	815	(Çetinkaya et al. 2002)
rpoB	C2700-F C3130-R	CGTATGAACATCGGCCAGGT TCCATTTCGCCGAAGCGCTG	446	(Khamis <i>et al.</i> 2004)
pld	pld-F pld-R2	ATAAGCGTAAGCAGGGAGCA ATCAGCGGTGATTGTCTTCCAGG	203	(Pacheco et al. 2007)

Table 1. Details of primer sequence used for detection of C. pseudotuberculosis.

were synthesised by Macrogen (South Korea) and the details are presented in Table1. The *16S rRNA*, *rpoB* and *pld* genes were amplified using ready-to-use master mixes for PCR (Ruby Taq Master®, Jena Bioscience, Thuringia, Germany). A total reaction volume of 25  $\mu$ l was used. Each reaction contains the following reagents: 12.5  $\mu$ l of master mix, 1  $\mu$ l of each forward and reverse primer (10 pmol), 2.5  $\mu$ l of template DNA and 8  $\mu$ l dH<sub>2</sub>O. PCR amplification was carried out in a Gene Amp PCR System 9700 Thermo Cycler.

The following PCR conditions (35 cycles) were used for amplification of *16S rRNA*, *rpoB* and *pld:* initial denaturation at 94 °C for 3 min, denaturation at 94 °C for 1 min, annealing at 56°C for 1 min, extension at 72 °C for 2 min and a final extension step at 72 °C for 7 min. PCR products were kept cold at 4°C until they were collected from the cycler. Amplified products were confirmed and separated by 1% (w/v) agarose gel electrophoresis made with TAE (1X) buffer stained with Prime Safe Dye (GeNet Bio, Korea). The fragment sizes were viewed under UV light.



Fig.1. Macroscopic appearance of affected lymph node from sheep and goats showing lesion of abscess in the center (1) and thick fibrous connective tissue as lamellate layer (2).

# **RESULTS AND DISCUSSION** Macroscopical observation

C. pseudotuberculosis is the causative agent of CLA in sheep and goats and causes economic loses in many countries. The aim of this study was to determine the infection rate of CLA in sheep and goats slaughtered at Duhok slaughterhouse and to correlate with the results of bacterial isolation with histophathological and molecular detection. From 1090 examined sheep and goats slaughtered at Duhok abattoir, 22 (2.1%) suspected CLA cases of mediastinal lymph nodes were collected from sheep and goats. Previously it has been reported that the prevalence rate of CLA among sheep and goats was variable and it was varied from country to county. Similar infection rates including 2.2%, 2.31% and 2.55% were recorded in Turkey, India and Iraq, receptively (Al-Badrawi and Habasha 2016, Cetinkaya et al. 2002, Kumar et al. 2013). However, the prevalence was reported to be 4.81% in Egypt (Mubarak et al. 1999) and it has also been shown to vary between 0.2% to 90.07% in different other studies (Al-Gaabary and Osman 2009, Al-Gaabary et al. 2010).

There was variation in the size of infected lymph nodes. Amongst all affecetd lymph nodes, mediastinal lymph node revealed marked congestion and oedematous changes. The present observations were in consonance to other studies as reported previously in goats and sheep affected with C. pseudotuberculosis (Sonawane et al. 2016, Singh et al. 2017). On careful examination extensive abscessation of mediastinal lymph nodes was prominent. The cut surface(s) of the mediastinal lymph node exhibited greenish yellow, thick semi solid ?uid mixed with pus arranged and trapped in concentrically lamellate layers of fibrous connective tissue (Fig. 1). Onion-like appearance was also found in some of the affected lymph nodes. The presence of onion-like layering of fibrous tissue was due to progressive stages of necrosis and capsule formation within abscess which given its pathogonomic lesion. (West and Bruere 2002). This

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Fig. 2. (A) Microscopic section of affected lymph node showing lesion of abscess (1), pyogenic membrane (2) with concentric fibrous layers separated as fibrous connective tissue (3); (B) Microscopic section of affected lymph node showing concentrically layer as onion appearance (1) and zone of pyogenic membrane (2) with infiltration of inflammatory cells and necrotic area (3); (C) Microscopic section of lymph node abscess revealing central caseo-necrotic core (1) surrounded by pyogenic membrane with infiltration of polymorphonuclear and mononuclear inflammatory cells (2) and fibrous capsule (3); (D) Microscopic section of lymph node showing abscess as a caseous necrosis (1) surrounded by pyogenic membrane with infiltration of polymorphonuclear and mononuclear inflammatory cells (2) (H & E 20X).

findings corroborated similarly to other studies (Ali *et al.* 2016, Mahmood *et al.* 2015) who observed the presence of similar lesion(s) following experimental inoculation of *C. pseudotuberculosis* in goats.

#### **Microscopical observation**

Histopathological lesions of affected lymph node(s) was characterized by the presence of central zone of abscess surrounded by pyogenic membrane with severe infiltration of polymorphonuclear and mononuclear inflammatory cells and concentric zones of connective tissue (Fig. 2A, B, C, D). In CLA the development of chronic lesions with multiple abscesses formation and lamellate layer are attributed to the ability of bacteria to escape the immune system and its intention to spread to many other organs. Virulence factor like phospholipase D

and mycolic acid (Baird and Fontaine 2007) determines its pathogenic ability, thus facilitates this organism to evade strong immune response from hosts. These findings were consistent to the previous reports (Sonawane *et al.* 2016, Singh *et al.* 2017) who stated the classical presentation of the histopathological feature of the pyogranulomatous in?ammation in lungs.

# Isolation and identification using standard microbiology methods

In addition to the post-mortem examination of infected lymph nodes, conventional laboratory methods have also been used for the diagnosis of suspected cases of CLA in sheep and goats (Dorella *et al.* 2006, Guimarães *et al.* 2011). The causative agent was identified as *C. pseudotuberculosis* based on colony characteristic after



Fig. 3. Colony characteristics, Gram stained smear and CAMP test of *C. pseudotuberculosis* isolates from sheep and goats.

A, non-hemolytic, dry and small white colonies after 24 h.B, narrow zone of â-hemolysis appeared 48 h incubation on %5 sheep blood agar. C, Gram-positive and club-shaped rods detected in stained smear with Gram stain. D, CAMP test of *C. pseudotuberculosis* (right) with *S. aureus* (center) shows inhibition of the effect of the beta-haemolysin produced by *S. aureus*.

24 to 48 h incubation, and also on the basis of morphological characteristics by Gram staining. On sheep blood agar small, dry and non-haemolytic white colonies were observed (Fig. 3A) and the colonies turned creamcolored colonies, with a narrow  $\beta$ -hemolysis zone, after further 48h of incubation (Fig. 3B). Gram-positive pleomorphic rods arranged as chinese letters were observed in Gram stained smear (Fig. 3C). The presumptive identification was confirmed by biochemical features. The isolates were catalase positive and they were able to hydrolyse urea. CAMP test showed inhibition of  $\beta$ -hemolysis by *Staphylococus aureus*as shown in Fig. 3D. The results of this study revealed that all caseous lymph nodes examined were found to be positive for *C. pseudotuberculosis*. It has been showed previously that the lymph nodes are the primary sites for replication of *C. pseudotuberculosis* and it has been identified from lymph nodes of sheep and goats (Çetinkaya *et al.* 2002, Ilhan 2013, Guerrero *et al.* 2018). Furthermore, *C. pseudotuberculosis* was isolated and identified in lymph nodes of mice in another study by Firdaus Jesse *et al.* (2013).

#### Molecular detection of C. pseudotuberculosis

The conventional bacteriological methods including colony morphology, biochemical test are usually not accurate due to the species variation within the genus *Corynebacterium* (Dorella *et al.* 2006). Therefore, fast and specific diagnostic tools are developed for early diagnosis of *C. pseudotuberculosis* especially from cases Histopathological and molecular studies on caseous lymphadenitis...



Fig. 4. Molecular detection of *C. pseudotuberculosis* isolates from sheep and goats.

1% agarose gel electrophoresis shows amplification of *pld*, *rpoB and 16S rRNA* genes specific for *C. pseudotuberculosis* from cases of CLA in sheep and goats. Lane 1: 100 bp DNA marker, Lane 2-4: amplification of 203 bp fragment of *pld*, Lane 5-7: amplification of 446 bp fragment of *rpoB*, lane 8-10 amplification of 815 bp fragment of *16S rRNA*.

of CLA by amplification of specific target genes including 16S rRNA, RNA polymerase  $\beta$ -subunit (*rpoB*) and phospholipase D (*pld*) genes (Çetinkaya *et al.* 2002, Khamis *et al.* 2004, Pacheco *et al.* 2007). In order to confirm the identification of the 22 suspected isolates by bacteriological methods, the isolates were further analysed using PCR. Detection was confirmed by amplification of the specific target genes of *C. pseudotuberculosis* including 16S *rRNA*, *rpoB* and *pld* respectively. All isolates were confirmed as *C. pseudotuberculosis* based on PCR amplification of 16S *rRNA* (815bp), *rpoB* (446 bp) and *pld* (203 bp) as shown in Fig. 4. Similar findings have also been confirmed in previous studies (Cetinkaya *et al.* 2002, Khamis *et al.* 2004, Hexian *et al.* 2018).

# CONCLUSION

Molecular and histopathological studies together could be used to identify *C. pseudotuberculosis* directly from cases of CLA from sheep and goats. Further studies including study of virulence genes and antimicrobial susceptibility together phylogenetic analyses are recommended to be done in the future for better understand the epidemiology of this pathogen in Duhok city.

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