



Traditional, histopathological and molecular diagnosis of sarcocystosis in slaughtered sheep in Al-Diwaniyah province, Iraq

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Abstract

The presented investigation was performed to identify *Sarcocystis* species, in an intermediate-host parasite with zoonotic importance, infection in slaughtered sheep identified by traditional, histological, and molecular methods in Al-Diwaniyah Province, Iraq. Here, 70 intercostal and esophageal muscle samples 10-40 gm/each were collected from slaughtered sheep. The samples were examined for detection of the Sarcocystic bradyzoites using microscopic, histological, *18S rRNA* gene-based polymerase chain reaction (PCR) and PCR-product-dependent gene sequencing methods. The results of the microscopic method revealed that 65 (93%) of the muscle samples contained bradyzoites. The histopathological picture showed the presence of two microsarcocysts with different morphological thin or thick wall sizes. The PCR demonstrated that 68/70 (97%) of the samples were positive for the occurrence of the parasite DNA. According to the NCBI-based websites, the Phylogenetic analysis revealed that the isolates from the current study were closely similar in their nucleotide sequences with isolates from Norway, Egypt, and Iran. The present data indicate high infection levels by *Sarcocystis* spp. in slaughtered sheep, which could bring alarm for public health importance.

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Introduction

As one of the Parasites of food-borne illnesses, *Sarcocystis* are intracellular protozoa. Until recently, 196 genuine species of *Sarcocystis* have been identified, and each of them is very selective regarding the types of hosts it can live on. Cysts are produced by parasites and may be seen in the striated muscles and the brains of pigs, cattle, and sheep (1-5). Eating raw or undercooked meat carrying sarcocystosis may spread infections of *S. hominis*, *S. heydorni*, and *S. suihominis*. Both *S. hominis* and *S. suihominis* have been designated as zoonotic risks by the European Food Safety Authority (6-12). Sheep may be infected with Sarcocystosis through drinking contaminated water or eating wild feed containing the parasite's sporocysts, leading to weight loss, abortion, early delivery, and even death (13-16). Most species of *Sarcocystis* need not one but two different hosts during their life cycles, with carnivores

servicing as the final hosts and herbivores acting as intermediate hosts. Only four parasitic species have been confirmed to infect sheep: the non-pathogenic parasitic species (macroscopic) *S. gigantea* and *S. medusiformis* transferred by felids, and the pathogenic parasitic species (microscopic) *S. tenella* and *S. arieticanis* spread by canids (17-21). *Sarcocystis* spp. Infects various animals, especially mammals, birds, and fish. The final host is responsible for the sexual phases of gametogony and sporogony. In contrast, the intermediate host is responsible for the asexual phases of merogony and cyst development (22,23). Sarcocystis infections in intermediate hosts are often asymptomatic, but severe instances have been observed, resulting in the host's fatigue, loss of appetite, diarrhea, loss of weight, muscular twitching, and in very exceptional instances, mortality. Muscle tissue cysts carrying bradyzoites are a characteristic feature of the degenerative alteration known as sarcocystosis, which occurs in the intermediate host.

Infections by *Sarcocystis ovicanis* or *Sarcocystis neurona* has been linked to rare cases of encephalitis in sheep and horses (24,25).

The presented investigation was performed to identify *Sarcocystis* species, in an intermediate-host parasite with zoonotic importance, infection in slaughtered sheep identified by traditional, histopathological, and molecular methods in Al-Diwaniyah Province, Iraq.

Materials and methods

Samples

The presented work collected 70 intercostal and esophageal muscle samples (10-40gm/each) from slaughtered sheep from April-December, 2022. The samples were cooled-box-transported to the Laboratory of Parasitology, College of Veterinary Medicine, University of Al-Qadisiya, Al-Diwaniyah City, Iraq.

Microscopic examination

The samples were examined for Sarcocystic bradyzoites and cysts using squeezing, trichoscopy, and solution analysis methods described by Al-Bayatee (26) and Al-Dulaimi (27).

18S rRNA gene-based PCR

18S rRNA gene-based PCR was performed using primers, Sar-F1: GCACTTGATGAATTCTGGCA and Sar-R1: CACCACCCATAGAATCAAG, followed by Stecher *et al.* (13). The QIAamp DNA Mini Kit (Qiagen, Germany), with steps from the kit itself was employed for the bradyzoite DNA extraction. A 1.5% agarose gel was placed under an electrophoresis process for 60mins at 100 volts and 80Amp. The gel, after that, was visualized under a UV-based imager. Stecher *et al.* (28) employed the methods in the current section.

18S rRNA-PCR product-based sequencing

The PCR-product-dependent gene sequencing method sends the PCR-purified products to Macrogen (Korea) for sequencing. The sequences were processed using MEGA X software and NCBI websites to complete the analyses and build a phylogenetic tree.

Results

Microscopic test

The results of the microscopic method revealed that 65 (93%) of the muscle samples contained bradyzoites (Figure 1).

Histopathological findings

The pathohistological picture showed the presence of two microsarcocysts with different morphological thin or thick wall sizes (Figure 2).

18S rRNA gene-based PCR findings

The PCR demonstrated that 68/70 (97%) of the samples were positive for the occurrence of the parasite DNA (Figure 3).



Figure 1: The bradyzoites of *Sarcocystis* spp. in sheep muscle tissue, 40X.

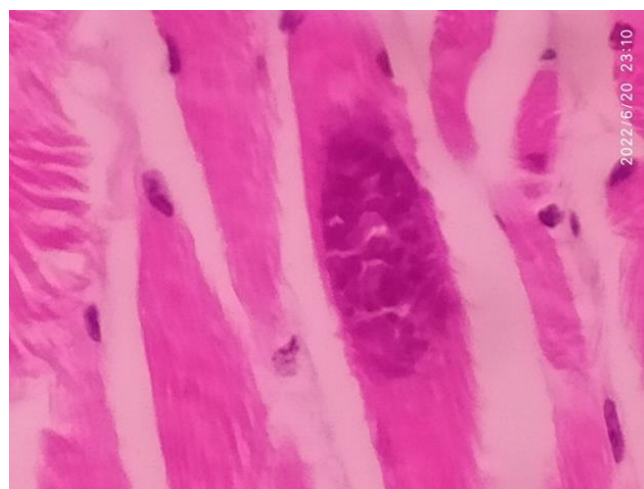


Figure 2: Encysted parasite in skeletal muscle showing internal bradyzoite nuclei separated by clear internal septations. H and E stain $\times 100$.

Findings of 18S rRNA-PCR product-based sequencing

According to the NCBI-based websites, the Phylogenetic analysis revealed that the isolates from the current study were closely similar in their nucleotide sequences with isolates from Norway, Egypt, and Iran (Table 1 and Figure 4).

Table 1: Homology sequence identity (%) in local *Sarcocystis* sp in sheep

Obtained Accession number	Identical to	GenBank Accession number	Country	Identity (%)	Host
OP835896	<i>Sarcocystis tenella</i>	MT560372	Iran	99.38	sheep
OP835897	<i>Sarcocystis tenella</i>	MT569891	Iran	100	sheep
OP835898	<i>Sarcocystis tenella</i>	MK420019	Norway	99.80	sheep
OP835899	<i>Sarcocystis tenella</i>	MG515221	Egypt	99.59	sheep
OP835900	<i>Sarcocystis tenella</i>	MT569891	Iran	100	sheep
OP835901	<i>Sarcocystis tenella</i>	MT569891	Iran	100	sheep
OP835902	<i>Sarcocystis tenella</i>	MT569891	Iran	100	sheep
OP835903	<i>Sarcocystis tenella</i>	MT569891	Iran	100	sheep
OP835904	<i>Sarcocystis tenella</i>	MT569891	Iran	99.59	sheep
OP835905	<i>Sarcocystis tenella</i>	MT569891	Iran	100	sheep

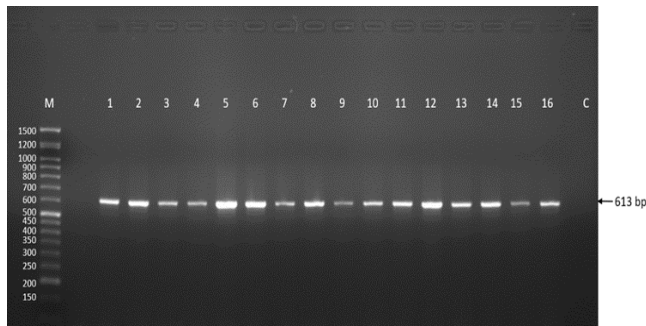


Figure 3: Image of agarose gel electrophoresis. It shows the *18S rRNA* gene-based PCR amplification of *Sarcocystis* spp. of sheep. 1-16: Positive samples. C: Negative control (No DNA). M: Molecular marker.

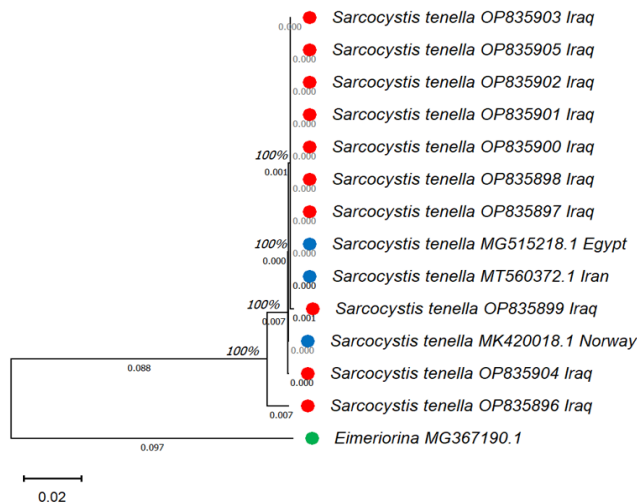


Figure 4: Phylogenetic tree of *Sarcocystis* spp in sheep based on the *18S rRNA* gene.

Discussion

Similar to the earlier findings from Baghdad 91.93% (26). this research found a prevalence of *Sarcocystis*

infection in sheep of 93%. And in Mongolia, the sheep have an infection rate of 97%, a 100% in Iran's sheep population, and 84 % in American sheep (29). Molecular methods, such as PCR amplification of species-specific 18s rRNA, are necessary for identifying species in this genus, which cannot be done only by microscopy. These results, at 97%, show that molecular diagnosis for sarcocystosis has a greater infection rate than traditional methods. Likewise, 90.78 % of the sheep samples in Karbala Province (30), Iraq, were infected with the parasite (31-33).

The *Sarcocystis* includes Sarcosporidia, known for their versatility in where they invade and grow inside the intermediate host. In addition, investigational transfer of the muscle cysts from sheep to dog puppies revealed the coccidian character of *Sarcocystis*, where the parasite was found at a higher incidence (34). *Sarcocysts*, only seen in the muscular parts of intermediate hosts, were characterized as evidence of a wide variety of host species and global dissemination. The current research demonstrated that natural infection rates in the tested hosts' skeletal muscles and organs varied. These findings are consistent with those found previously in Riyadh (35-37).

Alternatively, other nations reported lower levels. This research demonstrated a very varied rate of *Sarcocystis* distribution throughout several sheep organs. Oocyst exposure, changes in host immunity owing to abnormalities in the environmental and nutritional state, and parasite speciation are all possible causes for the observed variances in intensity among the tested hosts, as revealed by different studies (38). Other reports of detecting *Sarcocystis* spp. in different countries, such as Saudi Arabia, Tunisia, and China (39-41).

Conclusion

The present data indicate high infection levels by *Sarcocystis* spp. in slaughtered sheep, which could bring alarm for public health importance.

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Conflict of interests

The authors have not received any funding or benefits from industry, financing agency, or elsewhere to conduct this study.

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الطرق التقليدية، النسيجية المرضية والجزيئية لتشخيص الحويصلات الصنوبرية في الأغنام المذبوحة في محافظة الديوانية، العراق

هديل هادي جواد و غيداء عباس جاسم

فرع الأحياء المجهرية، كلية الطب البيطري، جامعة القادسية، الديوانية، العراق

الخلاصة

تم إجراء البحث المقدم هنا للتعرف على وجود خمج الحويصلات الصنوبرية، وهو طفيلي المضيف الوسيط وذو أهمية للأمراض حيوانية المصدر، في الأغنام المذبوحة التي تم تحديد الخمج بالطرق التقليدية والنسيجية والجزيئية في محافظة الديوانية، العراق. تم جمع ٧٠ عينة من العضلات ١٠-٤٠ جم / لكل منهما من الأغنام. تم فحص العينات لوجود الحويصلات بطيئة التكاثر باستخدام تفاعل البلمرة المتسلسل للجين الرابيوسومي وطرق دراسة تعاقب القواعد النيتروجينية لمنتجات تفاعل البلمرة المتسلسل. أظهرت نتائج الطريقة المجهرية أن ٦٥ (٩٣٪) من عينات العضلات احتوت على الحويصلات بطيئة التكاثر. أظهرت الصورة المرضية النسيجية وجود اثنين من الأوكياس المجهرية ذات حجم جدار رقيق أو سميك. أظهر تفاعل البلمرة المتسلسل أن ٧٠/٦٨ (٩٧٪) من العينات كانت إيجابية لوجود الحمض النووي للطفيلي في العينات المفحوصة. كشف تحليل تطور النسب أن العزلات من الدراسة الحالية كانت متشابهة إلى حد كبير في متواليات القواعد النيتروجينية مع عزلات من النرويج ومصر وإيران. تشير البيانات الحالية إلى مستويات عالية من الإصابة بطفيلي الحويصلات الصنوبرية. في الأغنام المذبوحة، الأمر الذي يمكن أن يندرج بأهمية الصحة العامة.