



Some Risk Factors Associated with Animal Brucellosis in Aswan Governorate, Egypt

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Abstract

A CROSS-SECTIONAL study was carried out during 2022 to evaluate the current status of bovine brucellosis at Aswan governorate as a border governorate and where an extensive process of live animal importation occurs through Abu Simbel quarantine. The present study estimated the seroprevalence of brucellosis among local and imported cows, identified the currently field circulating *Brucella* strains in Aswan governorate and described the most important risk factors associated with the dynamics of bovine brucellosis. A total of 932 cows including 145 imported cows and 787 local cows from different localities, 319 ewes and 581 does at Aswan governorate were serotested for brucellosis using Rose Bengal plate test (RBPT), and Buffered acidified antigen plate test (BAAPT) as screening tests and positive samples were confirmed by complement fixation test (CFT). The seroprevalence of brucellosis was estimated as 2 (1.4%) in imported cattle and 20 (2.5 %) in local cattle with overall prevalence of 22 (2.4%) in cattle, 4 (1.2%) ewes and 6 (1%) in does. Different clinical specimens including 10 cow's milk samples, 5 cow's aborted material and 5cow's manure samples were collected. Clinical samples were processed for bacterial isolation and PCR testing. A total eight *Brucella melitensis* biovar 3 and two *Brucella abortus* biovar1 isolates were recovered and identified on bacteriological and molecular bases. *Brucella melitensis* biovar 3 could be isolated from two manure samples of infected cows that suggests that manure could be a vehicle for indirect dissemination of infection in both animals and man and could be an important potential risk factor.

Keywords: Brucellosis, Manure, PCR, Risk factors, Sero-prevalence.

Introduction

Brucellosis is a highly contagious bacterial disease of sexually mature animals caused by different members of the genus *Brucella* [1]. Different types of *Brucellae* have their own specific and preferred natural hosts [2, 3, 4]. These hosts act as the most significant link maintaining the chain of infection of brucellosis through shedding of the pathogen by different routes achieving the persistence of the disease in infected regions. It is of importance to recognize that *Brucellae* are excreted significantly

from both aborted animals and cases of normal parturition of previously infected animals [5]. This leads to spreading the infection among herds especially in non-vaccinated cows.

As an intercellular pathogen, *Brucella* is a facultative intracellular pathogen that can multiply in macrophages avoiding the killing mechanisms by inhibition of phagosome lysosomes fusion [6]. Additionally, *Brucellae* have an efficient adaptation that prevents their recognition by the immune system [7].

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As the organism is capable of replicating inside macrophages [8, 9, 10], the multiplying Brucellae are then disseminated to other host cells infecting other organs. Brucella organisms have a predilection for placentas and fetal fluids in females and the testes of males [11]. Moreover, *Brucella melitensis*, *Brucella abortus* and *Brucella suis* are important cause of human illness representing one of the most serious neglected zoonotic diseases especially in developing countries [12, 13].

The epidemiology of bovine brucellosis is influenced by those factors that influence the maintenance of infection within both the infected animal and the surrounding environment. In addition, it is also influenced by the factors associated with disease transmission that lead to spread of infection among animals and between herds [14].

Brucella organisms have been confirmed to be excreted through the urine, milk, and abortion products of infected animals and up to 1×10^{13} organisms/gram have been associated with placenta and fluids as reported by [15]. Moreover, [16] found that Brucellae can be excreted in vaginal fluids associated with aborted fetuses, thereby contaminating feces and bedding of animals. Consequently, Brucellae can exist in the manure and environment of Brucella-infected animals and could be a vehicle for indirect dissemination and could be an important potential risk factor. Interestingly, [17] described an outbreak of *Brucella melitensis* in humans in Argentina that was attributed to contact with the manure of infected goats. In addition, high prevalence of Brucella-infection related with diminished proper manure removal, cleaning and disinfection procedures has been described by [18].

In animals, the main symptoms and sequelae suggesting brucellosis are late abortion, long inter-calving interval, infertility, premature birth and retained placenta [19]. However, following chronic infection, orchitis, swelling of the testes and hygromas can also be observed [20]. Clinical symptoms are not specific to brucellosis and must be confirmed by laboratory diagnosis.

Animal brucellosis mostly occurs due to the addition of infected animals to a susceptible herd, uncontrolled movement of animals especially in markets as well as mixing of different animal species [21, 22, 23, 24].

For the development of an effective control program for bovine brucellosis, it is essential firstly to recognize the potential risk factors that preserve the disease in both animals and the environment. Reservoirs that include Brucella-infected animals and contaminated environment especially the manure constitute the most important link of the chain of infection of brucellosis.

Therefore, the objectives of this study were focused on estimating the seroprevalence of brucellosis in imported and local cattle, ewes and does in Aswan governorate, identifying the currently field circulating strains of Brucella among cattle, describing the most important risk factors associated with brucellosis and discuss the role of Brucella infected animals and environmental reservoirs in the dynamics of animal brucellosis.

Material and Methods

Study areas and investigated animals

Different animal populations at different localities in Aswan governorate, Egypt, during 2022 including both imported and local cows, ewes and does were employed in this investigation (Figure 1). All animals were Brucella non-vaccinated and all populations were following the mixed breeding system especially with small ruminants.

Samples:

Ethical approval

The present study was approved by the "Ethical Committee", Beni-Suef University (BSU-IACUC/022-510), Egypt.

Blood sera

A total of 932 cows, 319 ewes and 581 does belonging to three districts (Kalabsha, Kom Ombo, and Sahari) at Aswan governorate, Figure (1) were selected and employed for serological diagnosis of brucellosis. Ten ml of blood were obtained from the jugular vein of each animal and left at room temperature to clot for serum separation. Blood sera were collected with plastic pipettes and transported in cold ice box to the laboratory stored at -20°C until tested.

Aborted material

Vaginal discharges and placentas of 5 aborted cows with retained placenta were collected. Cotyledons were separated and wrapped in a greaseproof paper, wrapped in strong plastic sacks, kept on ice and sent to the laboratory without delay for Brucella isolation.

Milk samples

Last streaks of milk were collected from 10 cows, and centrifuged at (6000 g) for 15 minutes. Cream and deposit mixture was used for Brucella isolation and PCR.

Manure samples

Five cow's manure samples were collected and processed from Brucella infected populations in private farms where Brucella infection was serologically confirmed in recently aborted cows. Samples were collected in sterilized plastic bottles directly from the rectum of each animal according to

the method described by [25]. Manure samples were then filtered using a syringe with a sterile cotton ball to remove solid debris twice, and the liquid was centrifuged at 3,000 x g for 30 min. Ten grams were added to 200 ml of tryptic soy broth, Becton Dickinson, 0.5% yeast extract, 5% horse serum, 1% glucose, and Brucella selective supplement), and incubated at 37 °C for 48 h in a shaking incubator. After centrifugation, 0.5 ml of the resulting pellet was used for Brucella isolation and PCR testing.

Serological examination

Blood sera were screened using Roe Bengal plate test (RBPT), and Buffered acidified antigen plate test (BAAPT) according to [26] and positive samples were confirmed by complement fixation test (CFT). Positive results were considered in samples that gave positive results in both screening and confirmatory tests.

Bacteriological examination

Milk cream and sediment mixture, vaginal discharges, placentas as well as pellets of manure samples were cultured on tryptose agar medium with selective antibiotic supplement (Oxoid), [27], according to [26] and the OIE [1] guidelines. Cultured plates were incubated in a CO₂ incubator with 10% CO₂. Vaginal discharges and the cream and deposit mixture of milk samples were spread on culture plates of the selective medium. Cotyledons were macerated and scrubbed onto the surface of the medium with and without antibiotics. The criteria used for Brucella identification included the requirement for additional atmospheric 10% CO₂, production of hydrogen sulfide gas, production of urease, growth on media containing the thionine and fuchsin, agglutination with monospecific antisera A, M and R and lysis by Tbilisi (Tb) and Izatnagar (Iz1) according to OIE [1] guidelines. Reference Brucella strains, *B. melitensis* Ether, *B. abortus* 544, and *Brucella suis* 1330 were supplied by the Central Veterinary Laboratory, Weybridge, Surrey KT15 3NB, UK and used as positive controls.

Molecular typing of Brucella isolates

DNA extraction

Milk, and manure samples were conveyed into new RNase-free Eppendorf tubes. Extraction was performed according to the instruction manual of Geneaid DNA extraction kit (New Taipei City, 22180 Taiwan, Cat. No. GS 100).

Conventional PCR

PCR amplification was carried out for molecular identification of *Brucella* spp. in DNA extracts from milk and manure at the genus level using primer sequences flanked the Immunodominant antigen, gene bp26 and Primer sequences flanked Erythritol catabolism, gene eryC according to [28] (Table 1).

Multiplex PCR

Multiplex PCR was performed for molecular identification of Brucella in DNA extracts from milk, and manure at the species level using five primers according to [29] (Table 1). The amplification was performed in Multigene thermal cycler (Labnet, USA).

The conventional and multiplex PCR thermal profiles were five minute at 94°C for initial heating followed by 30 cycles of 30 s at 94°C·45 s at 62°C, 1 min at 72°C, and final extension for 10 min at 72°C. The PCR amplicons were analyzed by running 7 µl of the PCR product in 1% agarose gel stained with ethidium bromide (0.5µg/mL). Gel documentation were used to visualize the gel under UV illumination. Control positive in conventional and multiplex PCR were illustrated in table (2)

Results and discussion

The seroprevalence of bovine brucellosis in the current study was estimated as 2 (1.4%) in imported cows and 20 (2.5 %) in local cows with an overall prevalence of 22 (2.4%) in cows, 4 (1.2%) in ewes and 6 (1%) in does based on RBPT and BAPAT as screening tests and CFT as a confirmatory test (Table 3). The results of this study are nearly comparable with other previous findings reported in different parts of Egypt. The prevalence of animal brucellosis in Egypt in the last few years ranged from 0.33% to 1.32 % [30]. Several reports have described the bovine brucellosis prevalence in Egypt with variable figures, 4.64% [31], 5.7% [32], 5.33% [33]. In sheep, the prevalence of brucellosis has been previously reported as 8.61% [34], 3.05% [35], 8.52% [31] and 8.2% [32]. Among goats, the prevalence of brucellosis was 11.3% [34], 3.64% [35], 8.51% [31] and 6.96% [32].

The difference between the prevalence obtained in this study and other previous studies could be due to the variation in management and breeding systems, the use of different diagnostic procedures, the presence of infected animals in the herds as sources of infection, introduction of infected animals, animals movement, and mixing of different animal's species.

An important factor responsible for the spreading of brucellosis is the uncontrolled importation from some brucellosis enzootic countries [22]. This was the common pattern in this study because Aswan is a border governorate through which many animals pass through the Abu Simbel quarantine where the introduction of live animals continuously occurs as a result of the urgent need for animal protein in Egypt. This is in addition to the illegal movement of animals across the open southern borders from neighboring enzootic countries including Sudan and Libya that is considered one of the major risk factors.

Moreover, movement of cattle between different animal's markets was observed in this study. This is considered as major risk factor and principal cause of failure of brucellosis eradication programs as previously discussed by [22, 24]. Moreover, [36] concluded that addition of animals in the herd is an important critical control point of brucellosis.

In this study mixed farming was observed as a criterion in the management system of animals in Aswan governorate where cattle were kept closely with small ruminants. Mixed farming of large and small ruminants has been reported to be a risk factor for *Brucella* transmission among different animal species as reported by [3, 37]. Cross-species infections frequently occur when different species are raised together. *Brucella melitensis* cross-infection from sheep and goats to cattle has been reported in southern Europe [38], the Middle East [39] and in Egypt [24].

The obtained results revealed isolation of 10 *Brucella* isolates. Among the 10 field *Brucella* isolates obtained from clinical specimens of cattle, eight were definitely *Brucella melitensis* biovar 3, Table 4 (80%) and two (20%) were bacteriologically identified as *Brucella abortus* biovar 1.

Brucella melitensis biovar 3 was previously described as the prevalent brucella type in Egypt as reported by [24, 39, 40, 41, 42].

Concerning *Brucella abortus* biovar 1, [40, 43] reported isolation of this type from 2 cows and 8 cows respectively in Egypt.

Isolation of *Brucella* organisms from aborted material (80%) is attributed to colonization of the reticuloendothelial system and genital organs causing chronic infection, reproductive disorders, abortion, stillbirth and infertility as explained by [44]. Moreover [45] concluded that pregnant cattle above five months of gestation are more likely susceptible to infection due to the preferential localization of *Brucella* in the uterus in which allantoic fluid factors such as erythritol stimulate the growth of *Brucella* organisms.

High association of *Brucella* seropositivity and abortion was observed in this study. To explain this association, it is important to recognize that *Brucella* infection is characterized by persistent infection in lymphoid tissues and inflammatory lesions in the reproductive tract of pregnant animals [44, 46]. Trophoblastic cells are the main target of *Brucella* infection during late gestation [44].

Retained placenta in this study was reported in all examined five aborted cows. Retention of the placenta can be explained on the basis of severe hemorrhagic necrotizing placentitis as described by [47].

Brucella abortus could be isolated from milk samples of 2 native cows. On the other hand, *Brucella melitensis* was isolated from milk samples of 2 native cows, 4 aborted materials of 4 imported cows and manure samples of 2 local cows. Isolation of *Brucella* organisms 4 (40%) from milk samples of cows coincides with our knowledge of considering the udder as an important portal of exit where the pathogen is shed. Cattle infected with *Brucella* spp. excrete high concentrations of the organism in their milk [48]. The recovery of *Brucella* from milk samples is of great public health significance as previously reported by [41].

Bacteriological examination revealed isolation of *Brucella melitensis* biovar 3 from two manure samples after exhaustive bacteriological work. Using the PCR assay the two manure samples revealed *Brucella melitensis* biovar 3 (Figure 2). Inappropriately, manure collected from animal farms is used as a fertilizer in most of the developing countries. Mitscherlich & Marth [49] reported the survival of *Brucella* in cow feces at room temperature for 122 days and 10 days in animal's environment. Remarkably, high humidity, low temperatures and lack of direct sunlight are the main factors leading to prolonged survival for *Brucellae* for months in water, aborted fetuses, placentas, liquid manure and hay as reported by [50]. The results obtained in this study suggest and add another mean of *Brucella* dissemination that complicates the infection chain of Brucellosis and may represent a potential risk for both animals and human health.

The results of conventional PCR confirmed the identification of *Brucella* in DNA extracts from milk, and manure on genus level using the genus-specific PCR with amplification of the fragment of 450 bp (Figure 2), as well as a fragment of 587 bp, specific for pathogenic *Brucella* strains of Erythritol catabolism, gene *eryC* (Figure 3).

Multiplex PCR in this study amplified the amplicons of 587, 1071, and 1682 bp indicating *Brucella melitensis* biovar 3, and the fragments of 587 bp, and 1682 bp sizes for *Brucella abortus* biovar 1, while *Brucella suis* reference strain has amplified the 272 bp, 587 bp, 1071 bp, and 1682 bp sizes (Figure 4). These findings agreed with those of [24, 41, 42] using similar primers.

Results shown in Figure (4) revealed no vaccine strains among the *Brucella* isolates. The obtained results confirmed the identification of both *Brucella melitensis* and *Brucella abortus*.

For preventive purpose, it seems it is essential to break the side of the epidemic triangle connecting the environment and the susceptible host. This can be done by highlighting the hygienic measures essential to limit the bacterial load in the environment and diminish the possibility of contact of animals with viable *Brucellae* in contaminated environments. The

important issues include the aborted materials which should be properly disposed as early as possible, the manure and the contaminated bedding that should be properly removed with removing of 20 cm. of the ground and replacing by a new layer mixed with lime and removing of shade to expose the area to the sun light as explained by [51].

Conclusion

This study revealed the enzootic occurrence of brucellosis in Aswan governorate, Egypt. Several risk factors are responsible for the transmission of brucellosis among animals that include importation of animals from enzootic countries, mixed farming and movement of animals and lack of vaccination in Aswan governorate in spite the enzootic pattern of the disease. Possible survival of *Brucella* organisms in manure of infected animals as proved in this study

suggests that manure could be a vehicle for indirect dissemination and could be an important potential risk factor.

Author contributions

All authors contributed in creating this article and approved the final manuscript.

Conflict of interest statement

Authors declare no conflict of interest

TABLE 1. Primers sequences, target genes, and amplicon sizes for conventional and Multiplex PCR.

Target gene	Primers sequences	PCR product (bp)
<i>Immunodominant antigen, gene bp26</i>	BMEI0535f GCG-CAT-TCT-TCG-GTT-ATG-AA	450
	BMEI0535r CGC-AGG-CGA-AAA-CAG-CTA-TAA	
<i>Erythritol catabolism, gene eryC</i>	BMEII0428f GCC-GCT-ATT-ATG-TGG-ACT-GG	587
	BMEII0428r AAT-GAC-TTC-ACG-GTC-GTT-CG	
<i>Glycosyltransferase, gene wboA</i>	BMEI0998f ATC-CTA-TTG-CCC-CGA-TAA-GG	1682
	BMEI0997r GCT-TCG-CAT-TTT-CAC-TGT-AGC	
<i>Outer membrane protein, gene omp31</i>	BMEII0843f TTT-ACA-CAG-GCA-ATC-CAG-CA	1071
	BMEII0844r GCG-TCC-AGT-TGT-TGT-TGA-TG	
<i>Erythritol catabolism, gene eryC (D-erythrulose-1-phosphate dehydrogenase)</i>	BMEII0428f GCC-GCT-ATT-ATG-TGG-ACT-GG	587
	BMEII0428r AAT-GAC-TTC-ACG-GTC-GTT-CG	
<i>ABC transporter binding protein</i>	BR0953f GGA-ACA-CTA-CGC-CAC-CTT-GT	272
	BR0953r GAT-GGA-GCA-AAC-GCT-GAA-G	
<i>Ribosomal protein S12, gene rpsL</i>	BMEI0752f CAG-GCA-AAC-CCT-CAG-AAG-C	218
	BMEI0752r GAT-GTG-GTA-ACG-CAC-ACC-AA	

TABLE 2. Lyophilized reference *Brucella* strains

Species	Biovar	Strain	ATCC	NCTC
<i>Brucella melitensis</i>	1	16M	23456	10094
		Rev.1		
<i>Brucella abortus</i>	3	Ether	23458	10509
		1		
<i>Brucella suis</i>	1	1330	23444	10316

ATCC¹: American Type Culture Collection, USANCTC²: National Collection of Type Cultures, UK.**TABLE 3. Seroprevalence of brucellosis at Aswan governorate**

Animals	Number of tested cows	Number of sero-positive cows*	Abortion
Imported cows	145	2 (1.4%)	10 (6.9 %)
Local cows	787	20 (2.5 %)	9 (1.14 %)
Total cows	932	22 (2.4%)	19 (2 %)
Ewes	319	4 (1.2%)	4 (1.2 %)
Does	581	6 (1%)	6 (1 %)

*In case of abortion, serum samples were collected three weeks after abortion

TABLE 4. *Brucella* isolation from different clinical samples at Aswan governorate

Samples	No. of samples	<i>Brucella</i> isolation	<i>Brucella abortus</i> 1	<i>Brucella melitensis</i> 3
Milk of native cows	10	4(40%)	2	2
Aborted materials of imported cows	5	4(80%)	-	4
Manure samples of native cows	5	2(40%)	-	2
Total		10	2(20%)	8(80 %)

Figure legends

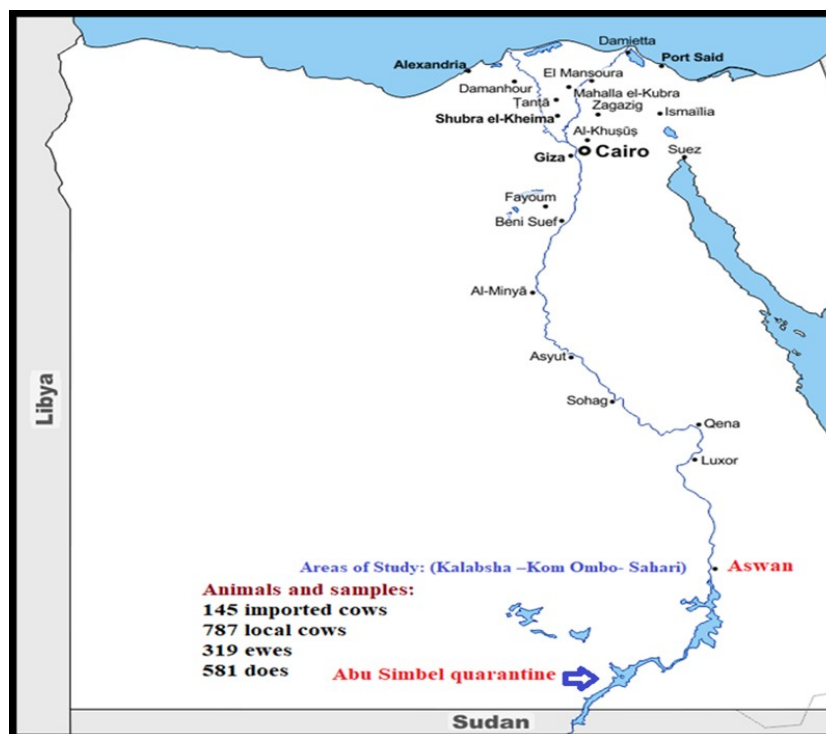


Fig. 1. Area of study, Animals and clinical samples



Fig. 2. Amplification of target gene immunodominant antigen, gene bp26 flanked the 450 bp. by conventional polymerase chain reaction (PCR).

Lane (1): 100bpDNA ladder.

Lane (2) *B. melitensis* reference strainLane (3:6) *Brucella* milk DNA extractLane (7, 8) *Brucella* manure DNA extract

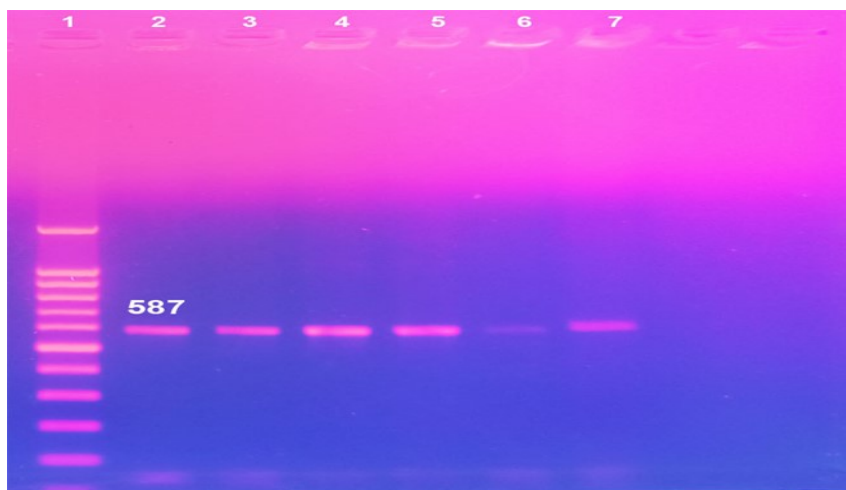


Fig. 3. Amplification of the set of primers specific for pathogenic Brucella strains flanked the 587 bp.

Lanes 1: 100bpDNA ladder

Lanes 2: 7, Brucella DNA extract.

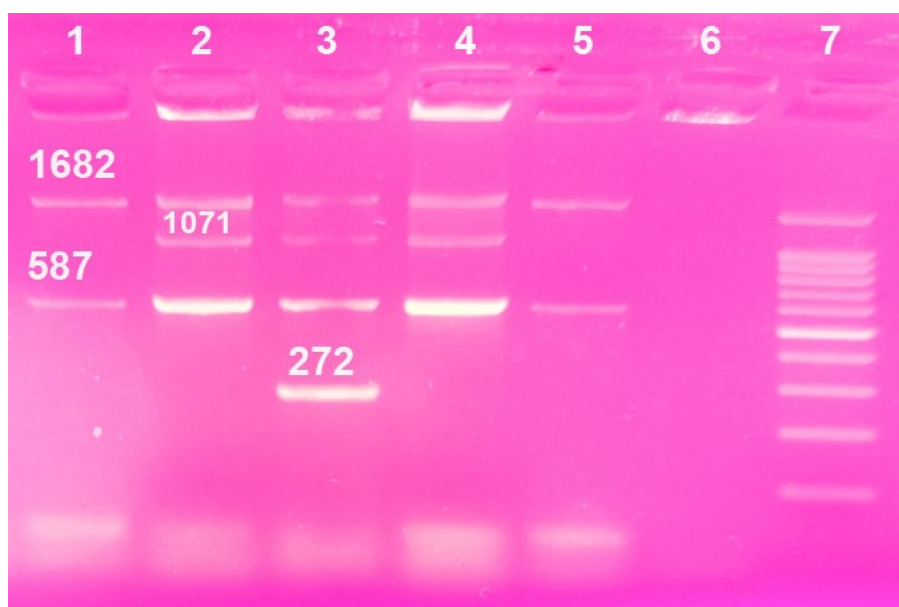


Fig. 4. Multiplex PCR of Brucella isolates

Lane (1) *B. abortus* 544 reference strain (587 - 1682 bp.)

Lane (2) *B. melitensis* reference strain Ether (587 - 1071 - 1682 bp.)

Lane (3) *B. suis* reference strain (272 - 587 - 1071 - 1682 bp.)

Lane (4) *B. melitensis* field strain (587 - 1071 - 1682 bp.)

Lane (5) *B. abortus* field strain (587 - 1682 bp.)

Lane (6) Control negative,

Lane (L) 100 bp marker.

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بعض عوامل الخطر المرتبطة بمرض البروسيلا في الحيوانات في محافظة أسوان، مصر

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أجريت هذه الدراسة خلال عام 2022 لتقييم الوضع الحالي لمرض البروسيلا في محافظة أسوان كمحافظة حدودية، حيث يتم استيراد الحيوانات الحية علي نطاق واسع من خلال محجر أبو سمبل. قامت الدراسة الحالية بتقدير معدل الانتشار المصلي لمرض البروسيلا بين الأبقار المحلية والمستوردة، وحددت سلالات البروسيلا المنتشرة حاليًا في محافظة أسوان، كما تم توصيف أهم عوامل الخطر المرتبطة بديناميكية انتشار مرض البروسيلا. تم اختبار 932 بقرة منها 145 بقرة مستوردة و787 بقرة محلية من مناطق مختلفة، و319 نعجة و581 نعجة في محافظة أسوان لمرض البروسيلا باستخدام اختبار روز البنغال (RBPT)، واختبار (BAAPT) تم تأكيد اختبارات الفحص والعينات الإيجابية عن طريق اختبار التثبيث التكميلي (CFT). تم تقدير معدل الانتشار المصلي لمرض البروسيلا (1.4%) في الأبقار المستوردة و20 (2.5%) في الأبقار المحلية، مع انتشار إجمالي قدره 22 (2.4%) في الأبقار و4 (1.2%) في النعاج و6 (1%) في النعاج. تم جمع عينات مختلفة اشتملت على 10 عينات من حليب البقر و5 عينات من المواد المجهضة من البقر و5 عينات من روث البقر. وتمت معالجة العينات لعزل البكتيريا واختبار PCR. تم عزل ثمانية عزلات من البروسيلا *melitensis biovar 3* وعزلتين من البروسيلا *abortus biovar 1* وتم تشخيصها على الأسس البكتريولوجية والجزيئية. تم عزل *Brucella melitensis biovar 3* من عينتين من روث الأبقار المصابة، مما يشير إلى أن الروث يمكن أن يكون وسيلة للانتشار غير المباشر للعدوى في كل من الحيوانات والإنسان ويمكن أن يكون عامل خطر محتملاً مهمًا.

الكلمات المفتاحية: البروسيلات، السماد، تفاعل البوليميراز المتسلسل، عوامل الخطر، الانتشار المصلي.