Detection of Brucella in infected bull testis by using PCR technique
الكشف عن جراثيم البروسيلا في خصية الثيران المصابة باستخدام تقنية تفاعل سلسلة البلمرة

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Abstract
Brucella infect male animals and cause orchitis and infertility, present study aimed to isolate brucella from testes of bull, 50 testes samples were collected then withdraw sample from epididymis after that stained with modified ziehl neelsen, samples were positive for stain were cultured on brucella selective media and tested with PCR. Results showed 6 samples were positive for modified ziehl neelsen, 4 samples were positive for culture and 6 samples were positive for PCR, from our research showed that PCR is more effective in diagnose Brucella infection.

الخلاصة
جراثيم البروسيلا تسبب التهاب الخصية والعمق في الثيران لذلك هدفت الدراسة إلى عزل جراثيم البروسيلا من خصية الثيران. تم جمع 50 عينة بعد ذلك سحب السائل المنوي من الأشعة باستخدام محاكيم طبية نبيذة ثم صبغت بصبغة الزيل نلسن المهورة وتم زرع وفحص العينات المعينة بصبغة الزيل نلسن المهورة باستخدام تقنية تفاعل سلسلة البلمرة. اظهرت النتائج 6 عينات كانت موجبة بصبغة الزيل نلسن المهورة و4 عينات موجبة للزرع وست منها كانت موجبة تقنية تفاعل سلسلة البلمرة. اظهرت نتائج البحث بحثا بأن تقنية تفاعل سلسلة البلمرة كانت الأفضل في الكشف عن الإصابة بجراثيم البروسيلا.

Introduction
Bovine brucellosis is usually caused by Brucella abortus, less frequently by B. melitensis, and occasionally by B. suis. Infection is widespread internationally. Several countries in Northern and Central Europe, Canada, Japan, Australia and New Zealand are believed to be free from the agent(1), (2). The disease is usually asymptomatic in nonpregnant females. Following infection with B.abortus or B.melitensis, pregnant adult females develop a placentitis usually resulting in abortion between the fifth and ninth month of pregnancy. In acute infections, the organism is present in most major body lymph nodes. Adult male cattle may develop orchitis and brucellosis may be a cause of infertility in male (3),(4). Hygromas, usually involving leg joints, are a common manifestation of brucellosis in some tropical countries and may be the only obvious indicator of infection; the hygroma fluid is often infected with Brucella(5). Presumptive evidence of Brucella is provided by the demonstration, by modified acid-fast staining of organisms, of Brucella morphology in abortion material or vaginal discharge, especially if supported by serological tests(6). The polymerase chain reaction methods provide additional means of detection. Whenever possible, Brucella spp. should be isolated using plain or selective media by culture from uterine discharges, aborted fetuses, udder secretions or selected tissues, such as lymph nodes and male and female reproductive organs. Species and biovars should be identified by phage lysis, and by cultural, biochemical and serological criteria. Polymerase chain reaction (PCR) can provide both a complementary and biotyping method based on specific genomic sequences(7). To the best of our knowledge there is no similar study in Iraq about orchitis in bull infected with Brucella.

Materials and Methods
A total of 50 suspected brucellosis testes samples were obtained from bulls carcasses with signs of orchitis(enlargement in testis, fever and hygroma in knee joints) were collected from Al-Samawa slaughter houses, sperms were withdraw directly from epididymis with sterile syringe and stained with Modified Ziehl Neelsen (MZN) , then cultured directly on brucella selective media then
certified *Brucella* growth with biochemical tests and monospecific antisera according to Alton(8), samples were positive for MZN were used for DNA extraction, was performed by using Promega DNA Isolation kit by using the primer set for IS711 genomic region of *Brucella spp.* as directed by Bricker & Hallin G (9) was commercially prepared and the sequences were as follows:

**F:** CAATCTCGAATTGGCCATCTCGAACGGTAT  
**R:** ATGTTATAGATGAGGTCGTCCGGCTGCTTGG

The PCR was performed in 50μl reaction mixture 1X Taq Buffer, 0.2mM dNTPs mixture, 1.5mM, MgCl2, 2.5U/μl Taq Polymerase, 4μM of each primer, 4μl of DNA extracted and 26.5 μl of DNase free deionized water. Each sample was tested in triplicate. The tubes containing the mixture were subjected to 35 cycles of amplification in a thermocycler. During each cycle the sample of DNA was denatured at 95°C for 35 seconds annealed at 64°C for 30 seconds, and extended at 72°C for 30 seconds. Prior to the cycling the mixture was subjected to incubation at 94°C for a period of 4 minutes. PCR product was then analyzed at 1.5% of agarose gel electrophoresis. The bands of *Brucella* spp. DNA were detected by using gel documentation system and gave an applicant size of 208bp.

### Results

Results showed 6 (12%) samples were positive for modified zielh Neelsen stain, *Brucella* organisms first recognized in smears obtained from epididymys stained with Modified Ziehl Neelsen stain, which appeared red clumps against a blue background (Fig.1), from results showed 4 isolates were obtained from epididymys. *Brucella* recognized on the basis of colonial morphology which appeared round translucent pale honey color on *Brucella* selective media, the results of biochemical test were positive for oxidase, catalase, nitrate reduction, H2S production and urease, negative for MR-VP, gelatinase, citrate utilization and indol production, all isolates were agglutinate with monospecific antisera for A. PCR was used to detect *Brucella* spp. in testes sample using the primers for IS711 genetic element and gave an applicant size of 208bp. The ladder used was 1500bp. PCR gave 6 positive result from 50 samples (Fig.2).
Fig. 4. Gel electrophoresis for PCR products where 208bp showed positive for *Brucella* spp.
Lane 1= marker, lane 2= *Brucella* isolate positive, lane 3= *Brucella* isolate positive,
lane 4= *Brucella* isolate positive, lane 5= *Brucella* isolate positive, lane 6= *Brucella* isolate positive,
lane 7= *Brucella* isolate positive.

Table (1). Positive samples for MZN, Culture and PCR assays.

<table>
<thead>
<tr>
<th></th>
<th>Test</th>
<th>+VE</th>
<th>%</th>
<th>-VE</th>
</tr>
</thead>
<tbody>
<tr>
<td>NO of samples</td>
<td>50</td>
<td>6</td>
<td>12</td>
<td>44</td>
</tr>
<tr>
<td>MZN</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Culture</td>
<td></td>
<td>4</td>
<td>8</td>
<td>46</td>
</tr>
<tr>
<td>PCR</td>
<td></td>
<td>6</td>
<td>11</td>
<td>44</td>
</tr>
<tr>
<td>Total</td>
<td></td>
<td>16</td>
<td>31</td>
<td></td>
</tr>
</tbody>
</table>

**Discussion**

The results showed that 6 samples were positive for MZN these results same with that reported by OIE, (1), was reported that *Brucella* are not truly acid-fast, but are resistant to decolorisation by weak acids and thus stain red by the Stamp’s modification of the Ziehl–Neelsen’s method. This is the usual procedure for the examination of smears of organs or biological fluids that have been previously fixed with heat or ethanol, and by this method, *Brucella* organisms stain red against a blue background. Only 4 samples were positive for culture and all isolates were *Brucella abortus*, this same that mentioned by (10), (11) whom concluded that biovar 1 is most frequently isolated from cattle, in countries where biovar prevalence has been studied. Outbreaks of bovine brucellosis are associated with abortion during the last trimester of gestation, and produces weak newborn calves, and infertility in cows and bulls. The primer pair used in this study succeeded in the amplification of a 208-bp fragment from epididymis samples were studied. In the meantime, the DNA extracted from milk harbor *Brucella*’s DNA results of PCR were the same as that obtained by Baily (12), (13) who certified that the PCR amplification contained a single pair of oligonucleotide primers designed to amplify a 223 bp product and reported that the assay was sensitive and specific for *B. melitensis* and *B. abortus*. PCR was used in the diagnosis of brucellosis and demonstrated it as an extremely specific, sensitive and easy and could become an usual diagnostic test for brucellosis. because then many studies described the PCR process for finding of the *Brucellae* in human and animals from special specimens. PCR process practical to human blood samples provide superior results than the conventional culture techniques for the diagnosis of together primary infection and relapses, as well as for focal complication of the disease (14), (15).
References:


