Neospora caninum Infection in Cattle in the Province of Kohgiluyeh and Boyer Ahmad, Southwest of Iran: Seroprevalence and Molecular Assessment

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1. Introduction

Neosporosis is an important disease in dogs and cattle with a global prevalence [1]. This disease is known as one of the main causes of abortion in cattle and neuromuscular paralysis in dogs [2]. The causative agent of neosporosis is an obligate intracellular protozoan and a member of the phylum Apicomplexa. One of the most important genus and species of which is Neospora caninum [3]. In cattle, the two main routes of Neospora infection are vertical transmission from mother to fetus, as well as transmission through ingestion of contaminated water or forage with oocysts excreted in dog feces [1]. Studies in different areas of the world show that a high percentage of aborted fetuses in cattle is due to infection with this parasite [4–6].

Neospora caninum causes miscarriage in cattle, and most miscarriages occur in the second trimester of pregnancy. The embryos are often autolyzed and may become dead or...
mummified in the uterus or even reabsorbed. Abortions occur throughout the year, both in the first and in subsequent deliveries [2]. Sometimes fetuses are born prematurely or seemingly healthy, without clinical signs [2].

Neosporosis is a common infection in livestock in Iran. According to a recent meta-analysis, the overall seroprevalence of bovine neosporosis in Iran ranges between 3.8 and 76.2% [7]. In a study by Sadrebazzaz et al. in Mashhad, East of Iran, 15.19% of cattle were found to be infected with *N. caninum*, and the rate of abortion in positive cattle was significant [8]. In another study in the same area, Razmi et al. reported that 46% of evaluated cattle are seropositive for neosporosis and that 85% of seropositive cattle had a history of abortions [9]. In a study in Kerman, southeast of Iran, on 285 bovine sera, *Neospora* infection was detected in 36% of the studied cattle [10]. In Ranjbar et al.’s study on aborted dairy cattle in Garmsar city, southeast of Tehran, 38.5% of studied animals were positive for *N. caninum* and there was a significant association between the prevalence of *Neospora* infection and the frequency of abortion in the studied cattle [11]. Rafati and Jafarian’s study on 100 aborted bovine embryos in Shahrekord in the southwestern part of Iran revealed that 11% of the samples are infected with *Neospora* parasite, by molecular method [12]. Ansari-Lari et al. reported a seroprevalence of 30% for *Neospora* infection in cattle in Shiraz, south of Iran [13].

A serological study of bovine neosporosis by Sengupta et al. in India revealed that 10% of the samples were infected with *N. caninum* and that there was a significant association between abortion and serum infection with *Neospora* [14]. A study in Argentina by Fort et al. on 4334 bovine serum showed that 302 (6.9%) of the cases were serologically positive for *N. caninum* infection [15].

Due to the high economic importance of this disease and also the existence of very different climates in different parts of Iran and the lack of sufficient information about this disease in livestock in different parts of the country, the present study is aimed at evaluating the status of *Neospora* infection in cattle in Boyer-Ahmad County in Kohgiluyeh and Boyer-Ahmad province, southwest of Iran, by molecular and serological methods.

2. Materials and Methods

2.1. Sample Collecting. In this cross-sectional study, 150 blood samples were obtained from cattle of Boyer-Ahmad County in Kohgiluyeh and Boyer-Ahmad province (Figure 1). The province is located in the southwest of Iran, where due to favorable weather conditions the prevalence of some of parasitic infection is substantial. Previous studies have reported a significant prevalence of helminth and protozoan infections in human as well as among livestock in the area [16–20].

The studied cattle in the current study were traditionally kept in stables and grazed from open fields and meadows. Information including age, sex, pregnancy status, and history of abortion was recorded in a predesigned datasheet. Of 150 cattle, 82 (54.7%) were male and 68 (45.3%) were female with no history of abortion. The studied cattle were between 1 and 8 years old, and the most common age group was the 5-year-old group, which included 23% of cases. The serum and buffy coat of the samples were separated and stored at -20°C until use. To do that, bloods were centrifuged (1000 g for 15 minutes). After centrifuge, at the junction of the plasma and red cells, there is a thin whitish or buffy coat layer. Serum, top layer, was first separated, and then, using a pipette, the buffy coat was collected.

2.2. Serological Evaluation of the Serum Samples

2.2.1. Preparation of Neospora Antigen. Vero cells were cultured in RPMI culture medium, containing 10% fetal calf serum, 100 IU/mL of penicillin, 100 μg/mL of streptomycin, 50 μg/mL of gentamicin, and 25 μg/mL of amphotericin. The *N. caninum* Nc1 isolate, provided by Razi Vaccine and Serum Research Institute of Shiraz Branch, Shiraz, Iran, was cultured in a Vero cell line. The culture was examined daily, and when 30% of the cell destruction by the parasite was observed, the Vero cells along with the tachyzoites were removed from the culture flask. The suspension was immediately washed 3 times, with PBS, and the supernatant was discarded. The tachyzoite pellet was washed in 2.3-ML of 37% formalin, and the concentration of formaldehyde was reduced to 6% with PBS washing solution and the obtained tachyzoite kept at 4°C, overnight for fixation. Then, formalin was removed by washing the sample with PBS. Alkaline buffer (BABS, pH: 8.7) was added to the tachyzoite suspension and that the final concentration of the tachyzoite was adjusted to 30,000 to 40,000 cells/μL.

2.2.2. Modified Agglutination Test (MAT) for Detection of Anti-Neospora IgG Antibodies. MAT was performed as previously described by Tavannae and Namavari [21]. Briefly, 95 μL saline was added to the first well of a 96 U-shaped microplate, and 50 μL was added to the rest of the wells. Then, 5 μL of cattle serum was added only to the first well of the plate, and from that well, 50 μL was transferred to the second well and continued in the same way until well 12 and the last 50 microliters were discarded. Then, 25 μL of 2ME solution was added to all wells and mixed thoroughly by shaking the plate for 4-5 minutes. Antigen (50 μL) diluted in borate buffer was added to each well, and the plate was shaken for 4-5 minutes as before. The plates were incubated in a wet chamber for 24 hours at room temperature. Finally, the plate was examined for the formation of agglutination under a loop microscope. Rabbit hyperimmune sera, raised against *Neospora* antigens, were used as positive control, and normal rabbit serum was used as the negative control.

2.3. Conventional PCR for Amplifying the 340 bp Fragment of the Neospora Nc-5 Gene. The genomic DNA from 127 cattle buffy coat samples were extracted, using a commercial Tissue Genomic DNA Extraction Kit (Favorgen Biotech Corp., Taiwan; FATGK001), following the manufacturer’s guidelines. A conventional PCR was performed to amplify a 340 bp fragment of the Nc-5 gene of *N. caninum*, using the NP6 forward (5′-CAGTCAACCTAGGTCTTCTTCT-3′)
and NP21 reverse (5′-GTGCCTCAATCTGTAAC-3′) primers.

A PCR reaction volume of 25 μL that consisted of 12.5 μL of 2x Taq PCR mix (Amplicon, Odense, Denmark), 0.5 μL of each primer (10 pmol/μL), 30 ng of template DNA, and 11.5 μL of ddH2O was prepared. The PCR temperature profile was one cycle at 95°C, 5 min, 30 cycles at 94°C, for 50 sec, 55°C for 30 sec, 72°C for 50 sec, and one cycle at 72°C, 4 min. In each run of the experiment, positive (N. caninum DNA) and negative (double-distilled water instead of template DNA) controls were included [22].

For separation of the DNA product, a 1.5% agarose gel in TAE solution was prepared and the PCR product was stained with GelRed nucleic acid gel stain for visualization under a gel documentation device.

2.4. Statistical Analysis. In this study, SPSS (ver. 20) software (SPSS, Chicago, IL, USA) was used to analyze the findings of the study. Chi-square test (χ²) was used to determine the association between the studied qualitative variables.

3. Results

Anti-Neospora antibodies were detected in sera of 49 out of 150 cattle corresponding to a seroprevalence rate of 32.7%. Even though the infection was more prevalent in five-year-old cattle, nevertheless, the differences between age and Neospora seropositivity were not statistically significant (p > 0.05). Table 1 shows the features of the cattle and relative seropositivity to Neospora in Kohgiluyeh and Boyer-Ahmad province, southwest of Iran.

Infection was more common in male cattle, yet the differences between sex and seropositivity to Neospora were not statistically significant (p > 0.05). PCR detected the Neospora DNA in theuffy coat of 26 out of 127 (20.47%) cattle (Figure 2).

The five-year-old cattle had the most (29%) positive molecular cases of Neospora infection, yet there was no statistically significant association between molecular positivity and age of the studied cattle (p > 0.05). Also, molecular positivity with Neospora was not associated with the sex of the evaluated cattle. Considering the agreement between the serological and molecular methods, 20 seropositive cases were PCR positive. The kappa coefficient test showed a moderate level of agreement (k = 0.73) between seroprevalence and molecular infection to Neospora in the animals.

4. Discussion

Neosporosis is a worldwide disease in which dogs and canines are definitive hosts in its life cycle [1]. Bovine neosporosis has been reported in different parts of the world, and it has been shown that 12-42% of cattle with a history of abortion have been infected with N. caninum [23].

The status of cattle infection with Neospora in many parts of Iran is unknown. Due to the importance of this parasite, it is worthwhile to determine the status of this parasitic infection in different species of animals, including cattle. These data are important for designing the control programs and to reduce the financial burden imposed by this parasite.

The present study was conducted, for the first time, in the southern regions of Iran to find out the status of neosporosis in cattle by serological and molecular approaches and this can be considered as the novelty of this study.

A total of 150 bovine blood samples were examined, using the MAT serological assay to detect N. caninum antibodies, and 49 cattle (32.7%) were found to be infected. Moreover, N. caninum DNA was detected in thebuffy coats of 26 out of 127 (20.47%) cattle, 20 of them were seropositive. The kappa coefficient test showed acceptable agreement between seroprevalence and molecular infection to Neospora in the studied animals. The overall seroprevalence of bovine neosporosis in Iran is reported to be 23.6% [4]. The findings which show a high prevalence of Neospora in the studied animals are in line with this report. In addition, in another study conducted in Fars province, which is located in the neighborhood of Kohgiluyeh and Boyer-Ahmad province, the prevalence of Neospora infection in cattle was reported to be 30% [13]. These studies indicate a high rate of infection with this parasite in cattle in Iran. A higher prevalence of Neospora (46%) has been reported from Mashhad in the East of Iran by Razmi et al. [9]. Also, higher prevalence for bovine neosporosis has been reported from other countries including Poland (56%) and Brazil (47.36%) [24, 25].

In the present study, Neospora infection was detected by the molecular method in a significant number of cattle.
Although this level of infection is consistent with the amount obtained by the serological method, the high rate of infection detected by molecular method in cattle cannot be simply justified. In most studies on Neospora infection in different hosts, the prevalence of Neospora by the molecular method has been much lower than that reported by the serological method. The same is true in our study, but the rate of infection by molecular method in our study is relatively high. In Duarte et al.’s study regarding the serological and molecular detection of N. caninum in human umbilical cord blood, the prevalence of Neospora infection was 24% by serological method and 1% by molecular one, using the umbilical cord blood [26]. However, this study utilized the umbilical cord blood for DNA extraction, and a low level of infection in a blood sample in neosporosis is not unexpected. Amdouni et al. in Tunisia also reported a 25% seroprevalence and a 12% molecular prevalence for Neospora infection in sheep, although the researchers performed the molecular experiment on the semen specimens [27]. A study in Shahrekord, southwestern Iran, revealed Neospora DNA in 11% of the aborted bovine embryos by molecular method [12]. A study by Hariri et al. detected Neospora antibodies in the sera of 6.7% of dogs in southern Iran, but none of them were positive by molecular method [28]. In the mentioned studies, samples other than buffalo coats have been used to identify the parasite DNA, while in the present study buffalo coat sample that seems to be a more suitable specimen has been used. This may partly explain the high molecular prevalence of Neospora detected in the present study.

In the current study, both molecular and serological methods were applied for the detection of Neospora infection in the studied cattle. For molecular evaluation, the Nc5 gene was targeted which is highly specific and excludes other species of toxoplasmidae family [29]. This can be considered as another strength of the present study. In addition, the presence of parasites’ DNA in the animal samples indicates the current state of Neospora infection in the animals. Furthermore, in the current study, serological method was used to assess the seroprevalence of neosporosis in the cattle. So far, several serological methods have been introduced to detect Neospora-specific antibodies in animal milk or serum, including ELISA, indirect fluorescent antibody tests (IFAT), and N. caninum modified agglutination test (MAT). These tests are commonly used in epidemiological studies, due to the fact that they detect both previous and current infections in animals and their positiveness indicates previous or recent infection with the parasite.

In the present study, the studied cattle grazed freely on pastures in rural areas where they were in constant and close contact with dogs, the definite host of the parasite. On the
other hand, keeping dogs in rural areas including the area where the current study was undertaken, particularly in families with livestock, is common. Therefore, it can be postulated that the high level of Neospora infection in the studied cattle is linked to their close contact with the dogs.

5. Conclusion

The findings of the present study showed a high rate of N. caninum infection in cattle in Boyer-Ahmad region in the southwest of Iran. This issue should be further investigated and the prevention and control of this parasite in livestock, due to the high financial burden of this parasite in the livestock industry, should be considered. Further studies on the prevalence of Neospora in other livestock in the region, as well as sequencing of PCR-positive samples and determining the genotype of the parasite in the region, are recommended.

Data Availability

The nominal and ordinal data used to support the findings of this study are available from the corresponding author upon request.

Ethical Approval

The study was approved by the Ethics Committee of Shiraz University of Medical Sciences (SUMS) (Ref. No. IR.sums.med.rec.1397.574).

Conflicts of Interest

The authors declare that they have no conflicts of interest.

Authors’ Contributions

BS and MN designed the study. AD, NA, SS, SZ, AM, and MZ were involved in data collection and performing the experiments. All authors read and approved the final version of the manuscript.

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