

Divergent Diagnostic Approaches in Investigating Brucellosis: Unveiling Clinical Signs and Infection Rates

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Abstract

Brucellosis, a zoonotic infectious disease caused by bacteria of the Brucella species, remains a significant global public health concern, impacting both humans and animals. To arrive at a thorough comprehension of the topic, an extensive evaluation was conducted, which involved examining 60 carefully obtained serum specimens from various demographic groups. These groups comprised veterinary students, those in charge of animal care, paraveterinarians, licensed veterinary technicians and experts in the complex field of synthetic animal conception. The study sought to provide a nuanced view on the number of cases and possible factors of brucellosis among the veterinary healthcare industry in the designated geographic region by deliberately included such a varied array of individuals. The study examined sera sample for brucellosis using various methods, including the "serum agglutination test (SAT), Immunoglobulin G (IgG) and Import General Manifest (IGM) indirect ELISA, Rose Bengal plate test (RBPT) and polymerase chain reaction (PCR)". Demographic information such as clinical symptoms, age, risk factors and sex was collected through a structured questionnaire. Intermittent fever (71.62%) was the most common clinical sign observed, along with discomfort throughout the body. The guidelines for early identification, therapy and sickness follow-up highlight the need of a high degree of suspicion, perception and a multimodal testing method.

Keywords: Brucellosis, Animals, Risk Factors, Artificial Insemination.

INTRODUCTION

Human infections are zoonotic and among the most common zoonosis worldwide is brucellosis, which is brought by bacteria from a species of Brucella (1). Of twelve forms of Brucella is that are currently identified, three were reported as causing severe illness and impairment in humans: Brucella Suis, Brucella melitensis and the bacteria Brucella abortus (2). Bravecellosis is a difficult infection that affects humans and cattle. Its prototypical clinical presentations, diagnostic limits, protracted treatment courses and recurring fever episodes make it such. Numerous symptoms, including fever, perspiration, malaise, anorexia, headache, the joint and vertebral pain, might be indicative of the condition (3). The two major ways that humans might become infected from diseased animals are through contact with them and through eating unsterilized milk and meat products. Blood transfusions, bone marrow transplants, intimate relationships, transplacental dissemination, exposure to the mother's meteorites after birth and breast breeding are some of the ways that infections can spread from person to person (4). Despite the availability of cultural, serological and molecular diagnostic methods, the nonspecific clinical characteristics of brucellosis pose a problem for doctors in terms of prompt and accurate identification of human infection. Sluggish blood culture development had the difficulty of serodiagnosis (5). Although it is a tedious and risky procedure for lab personnel, the isolation of bacteria from the patient's blood, stem cells, or other kinds of tissue is the gold standard and confirmed diagnosis. Serological testing is recommended to prevent false positive and negative findings as well as to quickly reduce the time needed for numerous tests by amplifying DNA targets using various PCR techniques (6). Increased risk of spreading the illness to animal handlers, vets, veterinary workers, vaccinators and school children is associated with high the



infection frequency in mammals. Despite the advances in medical research, certain challenges persist in the field of epidemiology. A significant barrier to comprehensive data collection stems from gaps in knowledge, compounded by insufficient laboratory facilities and the prevalence of misdiagnoses arising from overlapping clinical spectra (7). These impediments have led to a scarcity of information crucial for understanding and managing the epidemiology of various illnesses. In light of these limitations, the primary objective of the current investigation is to employ advanced multimodal differentiated diagnostic techniques. The approach aims to determine the prevalence of cases, identify associated signs along with symptoms and assess potential hazards associated with the targeted illness. The information gathered is crucial for planners creating animal control strategies, as well as for epidemiologists looking for a more realistic representation of disease trends (8). Furthermore, this understanding can improve diagnosis accuracy, allowing for early therapy commencement and ongoing patient monitoring, which will be beneficial to healthcare practitioners. Through addressing these important gaps, this research hopes to make a significant contribution to the care of humans and animals equally.

The study (9) created a GP-pharmacist collaborative paradigm, the study sought to explore the convergence and conflicting views and perspectives of common community pharmacists (CPs) toward and their professed hurdles of partnership. The study (10) examined container-casualty rates varying from the total number of fatalities reached 12,784. The elderly and those with comorbidities have the greatest death rates from the condition, whereas children appear to be the least afflicted demographic. The study (11) examined when someone coughs, sneezes, comes into touch with their lips or eyes, or comes into contact with infected hard surfaces. SARS-CoV-2 propagation is accelerated by close human contact. The study (12) evaluated the relationship with monthly cultures and NGS in subsequent knee pasty. Every patient having a second shoulder surgery was recruited systematically. The study (13) evaluated shed light on these topics by contacting organizations and networks that are associated with arrhythmias. A set of fifteen questions on CIED viruses was given to members of seven arrhythmia societies throughout the globe. The work (14) determined the ideal read values for identifying common infections and to conduct a thorough examination of micro biome next-generation sequencing for the etiological diagnosis of patients with septic conditions. The study (15) emphasized the finding that Candida albicans and Candida glabrata take distinct routes that lead to a comparable result. It draws attention to the gaps in our understanding of some of the particular processes underpinning the disease processes of the parasite, which call for more investigation. The study (16) concentrated on the clinical and sociological information from 533 HPS cases that the National Reference Laboratory for Hantavirus verified between 2009 and 2017. The study (17) evaluated when various adherence habits affected the long-term efficacy of Cyclic Redundancy Check (CRC) screening for a screening colonoscopy at 10-year intervals and yearly vomit immunochemical screening. The study (18) examined molecular characteristics between patient-matched diagnostic and reverted tumours and to look into the clinical effects of relapsed medulloblastoma. DNA methylation array and nextgeneration sequencing were used to examine a matched molecular cohort that was mainly unbiased. The study (19) combined the most current data that was made available for every step of the COVID-19 result continuum population, experience, infection diagnosis, death and hospitalization. The primary clinical characteristics of COVID-19 and the possible involvement of macrobiotic in COVID-19 are discussed in the following summary [20].

METHODOLOGY

Brucellosis, resulting from the infection by different species of the genus Brucella, is primarily observed in animals; however, transmission to humans can occur through contact with either infected animals or their derivatives.



Data collection

The study included 60 patient samples that were obtained from nearby hospitals and 30 control samples that came from functioning properly, unaffected volunteers. Each subject had their written consent acquired after being briefed about the research methods [21].

Selection criteria

- A medical background involving contact with animals, enduring symptoms like prolonged fever, sweating, chills and persistent joint/body/back discomfort for over two weeks, along with a notable antibody titre (>1:160) in serological agglutination testing (SAT).
- If clinical symptoms indicative of brucellosis persist for over two weeks, along with a documented history of animal association, the identification of anti-Brucella antibodies requires the use of at least two serological tests.
- Regardless of seropositive status, amplification by PCR of DNA is shown when combined with an illness suggesting a connection to animals and the existence of enduring symptoms suggestive of brucellosis that continue for longer than two weeks.

Sample Preparation

In the course of a brucellosis study conducted in diverse districts of Karnataka, a collective of 60 blood samples was collected from participants, encompassing par veterinarians, individuals engaged in artificial insemination, veterinary students, veterinarians and animal handlers. The individuals voluntarily gave permission to share personal information, including age, location, sex, employment, history of animal associations and ailments. The spoken and printed consent were obtained for the compilation of 5 ml of blood using anticoagulant-free vacationers.

Calculation of percent positivity (PP)

The colour was created through the utilization of "*o-phenylenediamine dihydrochloride* (OPD) as the *chromogen and hydrogen peroxide* (H_2O_2) " as the substrate. IELISA was used for measuring optical densities (OD) at 492 nm and the accompanying formula-1 was used to determine the (PP) factors.

$$PP = \frac{Average \ OD \ values \ of \ test \ sera}{Median \ OD \ value \ of \ strong \ positive \ control \ sera} \times 100 \tag{1}$$

PCR Amplification

MedCalc software was used for contrasting the PP values acquired from iELISA with 2-ME-SAT and SAT titles. The positive cut-off, determined as $\geq 50\%$ PP, was particular base on the greatest compassion and specificity values obtain from the ROC curve. The sera samples underwent DNA extraction using the QIAamp DNA kit in accordance with the manufacturer commands. (F):"B5 (R): CGCGCTTGCCTTTCAGGTCTG and TGGCTCGGTTGCCAATATCAA." A 25 μl effect quantity for the type-exact PCR contained two X Go Taq green masters mixes, 0.2 μM of B4 and B5 primer and around 30 ng of serum DNA. In a thermal cycler, there were 35 cycles consisting of a minute at 93 °C, one minute at 60 °C and 45 sec at 72 °C.

Evaluation of Positivity

The serum samples underwent initial testing using the rapid screening test, RBPT. Samples testing positive in RBPT were subjected to SAT using twofold consecutive dilutions, opening from 1:10 to 1:1280 strength, following the Weybridge method. The endpoint titre for human brucellosis was determined as the maximum strength of the serum exhibiting 50 percent agglutination. A titre of 1:160 (320 IU/ml) and higher was deemed



positive. Anti-Brucella IgM and IgG antibodies in sera sample be screen using laboratory-consistent iELISA protocol.

IELISA human restorative

The iELISA, human convalescent sera were employed as IgM and IgG favourable restrictions have demonstrated positivity in RBPT. To establish downbeat organize in ELISA, a serum example obtained from a healthy donor, confirmed negative for brucellosis through all relevant tests and it was employed. Titrating the target antigen beside 1:100 as well as 1:200 concentrations of both power negative and positive sera sample in a jamming cushion, the ideal concentration of antigen for IELISA was found.

Statistical Analysis

The Chi-square test was utilized to evaluate the relationship between risk indicators and seropositivity across the specified variables: education, region, sex, age and occupation. The analysis was conducted using the SPSS 22 program, with a defined threshold of statistical significance set at a p-value of ≤ 0.05 .

RESULT

The current research examined 60 sera samples obtained from individuals in veterinary healthcare, assessing them for brucellosis antibodies through various tests. Out of these, 35 (58.3%) were males, while 15 (25%) were females. The people in the group ranged in age from 21 to 69. The age category ranging from 31 to 40 accounted for the bulk of samples (16%), which was followed by those of 41 to 50 (35%). Moreover, 16% of the participants belonged to the age bracket of 21 to 30. Older age ranges (51–60 and above 60). Significantly, seroprevalences of the illness were higher in the age's ranges of 21–30 (14.28%), 41–50 (65%) and 31–40 (84%) (Table (1)).

Risk factor	Features	Number of	Number of	Number of	χ^2	p-value
		sample	Positive	negative		
Gender	Female	15	3 (20%)	12 (80%)	4.71	0.029*
	Male	35	5 (14.28%)	(85.71%)		
Age in years	21-30	7	1 (14.28%)	6 (85.71%)	3.4	0.524
	31-40	25	4 (16%)	21 (84%)	1	
	41-50	20	7 (35%)	13 (65%)		
	>60	8	1 (12.5%)	7 (87.5%)		
Education	Post graduates	18	7 (38.9%)	11 (61.11%)	2.85	
	Graduates	23	3 (13.04%)	20 (86.95%)		
	higher secondary	5	0 (0%)	5 (100%)		0.579
	Junior to senior high school	10	2 (20)	8 (80%)		
	Un educated	4	1 (25)	3 (75%)		
Profession	Veterinary students	9	4 (44.4%)	5 (55.6%)		
	Artificial inseminators	12	3 (25%)	9 (75%)		

 Table (1). Outcomes of medical and serological identification of hypodermis's in 60 Brucellosis (Source: Author)



	Para-	4	2 (50%)	2 (50%)		<0.0001*
	veterinarians					
	Veterinarians	25	7 (28%)	18 (72%)		
	Animal	10	3 (30%)	7 (70%)		
	handlers					
Region	Rural	37	7 (18.91)	30 (81.08%)	4.14	
	Urban	23	3 (13.04%)	20 (86.95%)		0.042*

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In various high-risk categories, veterinarians constituted 4 (44.4%), animal handlers accounted for 3 (30%), veterinary students represented 7 (28%), par veterinarian staff comprised 2 (50%) and individuals involved in artificial insemination constituted 3 (25%). Notably, higher prevalence rates of brucellosis were observed among animal handlers (70%) and par veterinarians (16.32%) in comparison to veterinarians (72%) along with those engaged in artificial insemination (75%).

The statistical analysis revealed that individuals involved and veterinary students faced a lower risk of infection (p-value < 0.05) compared to para-veterinarians, animal handlers and veterinarians. Table (2) depicts the most common observed symptom was a fever that fluctuated (71.62%), epididymo-orchitis in (12.16%) of cases which was followed by reproductive problems in (8.10%) of patients and spondyloarthropathy in (52.70%) of patients.

Medical grievances	No. of cases		
Sporadic fever	53 (71.62%)		
Body aches	33 (44.59%)		
Weight loss	23 (31.08%)		
Abdominal pain	13 (17.56%)		
Depression	4 (5.40%)		
Infertility			
Joint pain	39 (52.70%)		
Sweating	31 (41.89%)		
Head ache	16 (21.62%)		
Pain in testicles	9 (12.16%)		

Table (2). Number of Medical complaints (Source: Author)

DISCUSSION

The worldwide public health and food security are at danger from brucellosis, which is recognized as an illness of the workplace. In a number of nations, counting India, the true incidence for human brucellosis is uncertain. During the past 25 years, there has been a dynamic evolution in the epidemiological landscape of human brucellosis, with new focus areas emerging and re-emerging constantly. Many livestock farms, hygienic conditions, social pressures and politics are some of the elements that are blamed for this change. Brucellosis has been identified as a considerable contributor to morbidity related to travel. Human brucellosis can cause lifethreatening symptoms, however it manifests as a feverish disease with minimal mortality. Comparing veterinarians (28%) and artificial inseminators (25%), the prevalence of brucellosis is much greater among paraveterinarians (50%) and animal handlers (30%). Among those who provide care for animals, there was a significant correlation with brucellosis (p < 0.0001). Brucellosis in humans has been documented in cases of pyrexia of unknown origin (PUO), as well as among animal handlers, veterinarians and slaughterhouse workers in India. A large number of the prevalence of infection studies, hospital-based observation reports and instances



has been released. It has been discovered that gender is an individually important variable linked to a sickness known as a disease that only affects men.

CONCLUSION

The transfer of brucellosis from animals to humans is a well-established occurrence, manifesting through direct contact and the ingestion of tainted food. The endemic prevalence of the disease, coupled with the diverse and mixed farming practices in livestock, poses a significant risk to the health of individuals involved in animal care. Therefore, by implementing an extensive initiative to employ appropriate protective measures when dealing with infected animals can diminish the occurrence of the disease among animal handlers and veterinary healthcare professionals. To successfully control the condition and reduce the significant mortality related to it, it is imperative to create a well-defined test procedure to distinguish between acute and chronic infections in groups at risk, as well as to conduct frequent serological follow-ups. To emphasize the significance of prevention, it is imperative to stress the screening of relatives in endemic regions for acute cases of brucellosis.

REFERENCE

- Yagupsky, P., Morata, P., & Colmenero, J. D. (2019). Laboratory diagnosis of human brucellosis. Clinical microbiology reviews, 33(1), 10-1128. DOI: <u>https://doi.org/10.1128/cmr.00073-1</u>
- [2] Muñoz, P. M., Mick, V., Sacchini, L., Janowicz, A., de Miguel, M. J., Cherfa, M. A., ... & Garofolo, G. (2019). Phylogeography and epidemiology of Brucella suis biovar 2 in wildlife and domestic swine. Veterinary microbiology, 233, 68-77. DOI: <u>https://doi.org/10.1016/j.vetmic.2019.04.025</u>
- [3] Mishra, A. R., Rani, P., Krishankumar, R., Ravichandran, K. S., & Kar, S. (2021). An extended fuzzy decision-making framework using hesitant fuzzy sets for the drug selection to treat the mild symptoms of Coronavirus Disease 2019 (COVID-19). Applied soft computing, 103, 107155. DOI: <u>https://doi.org/10.1016/j.asoc.2021.107155</u>
- [4] Yadav, D. D., Singh, N., Sreedharanunni, S., Hira, J. K., Chhabra, S., Trehan, A., ... & Das, R. (2021). Extreme Genotype/Phenotype Heterogeneity of Double Heterozygous Sickle β-Thalassemia in a Family: Implications in Antenatal Diagnosis. Indian Journal of Hematology and Blood Transfusion, 37, 689-691. DOI: 10.1007/s12288-021-01412-1
- [5] Sridhar, S., Situ, J., Cai, J. P., Yip, C. C. Y., Wu, S., Zhang, A. J. X., ... & Yuen, K. Y. (2021). Multimodal investigation of rat hepatitis E virus antigenicity: Implications for infection, diagnostics, and vaccine efficacy. Journal of Hepatology, 74(6), 1315-1324. DOI: <u>https://doi.org/10.1016/j.jhep.2020.12.028</u>
- [6] Richardson, S., Hill, R. M., Kui, C., Lindsey, J. C., Grabovksa, Y., Keeling, C., ... & Clifford, S. C. (2022). Emergence and maintenance of actionable genetic drivers at medulloblastoma relapse. Neuro-oncology, 24(1), 153-165. DOI: https://doi.org/10.1093/neuonc/noab178
- [7] Hu, T. H., Rosli, N., Mohamad, D. S., Kadir, K. A., Ching, Z. H., Chai, Y. H., ... & Singh, B. (2021). A comparison of the clinical, laboratory and epidemiological features of two divergent subpopulations of Plasmodium knowlesi. Scientific reports, 11(1), 20117. DOI: <u>https://doi.org/10.1038/s41598-021-99644-8</u>
- [8] Forsea, A. M. (2020). Melanoma epidemiology and early detection in Europe: diversity and disparities. Dermatology practical & conceptual, 10(3). DOI: <u>https://doi.org/10.5826%2Fdpc.1003a33</u>
- [9] Saha, S. K., Kong, D., Thursky, K., & Mazza, D. (2021). Divergent and convergent attitudes and views of general practitioners and community pharmacists to collaboratively implement antimicrobial stewardship programs in Australia: a nationwide study. Antibiotics, 10(1), 47. DOI: <u>https://doi.org/10.3390/antibiotics10010047</u>
- [10] Abduljalil, J. M., & Abduljalil, B. M. (2020). Epidemiology, genome, and clinical features of the pandemic SARS-CoV-2: a recent view. New microbes and new infections, 35, 100672. DOI: <u>https://doi.org/10.1016/j.nmni.2020.100672</u>
- [11] Machhi, J., Herskovitz, J., Senan, A.M., Dutta, D., Nath, B., Oleynikov, M.D., Blomberg, W.R., Meigs, D.D., Hasan, M., Patel, M. and Kline, P., 2020. The natural history, pathobiology, and clinical manifestations of SARS-CoV-2 infections. Journal of Neuroimmune Pharmacology, 15, pp.359-386. DOI: <u>https://doi.org/10.1007/s11481-020-09944-5</u>
- [12] Namdari, S., Nicholson, T., Abboud, J., Lazarus, M., Ramsey, M. L., Williams, G., & Parvizi, J. (2019). Comparative study of cultures and next-generation sequencing in the diagnosis of shoulder prosthetic joint infections. Journal of shoulder and elbow surgery, 28(1), 1-8. DOI: <u>https://doi.org/10.1016/j.jse.2018.08.048</u>
- [13] Traykov, V., Bongiorni, M. G., Boriani, G., Burri, H., Costa, R., Dagres, N., ... & Blomström-Lundqvist, C. (2019). Clinical practice and implementation of guidelines for the prevention, diagnosis and management of cardiac



implantable electronic device infections: results of a worldwide survey under the auspices of the European Heart Rhythm Association. EP Europace, 21(8), 1270-1279. DOI: <u>https://doi.org/10.1093/europace/euz137</u>

- [14] Ren, D., Ren, C., Yao, R., Zhang, L., Liang, X., Li, G., ... & Gui, S. (2021). The microbiological diagnostic performance of metagenomic next-generation sequencing in patients with sepsis. BMC Infectious Diseases, 21, 1-9. DOI: <u>https://doi.org/10.1186/s12879-021-06934-7</u>
- [15] Galocha, M., Pais, P., Cavalheiro, M., Pereira, D., Viana, R., & Teixeira, M. C. (2019). Divergent Approaches to Virulence in C. albicans and C. glabrata: Two Sides of the Same Coin. International journal of molecular sciences, 20(9), 2345. DOI: <u>https://doi.org/10.3390/ijms20092345</u>
- [16] Alonso, D. O., Iglesias, A., Coelho, R., Periolo, N., Bruno, A., Córdoba, M. T., ... & Martínez, V. P. (2019). Epidemiological description, case-fatality rate, and trends of Hantavirus Pulmonary Syndrome: 9 years of surveillance in Argentina. Journal of Medical Virology, 91(7), 1173-1181. DOI: <u>https://doi.org/10.1002/jmv.25446</u>
- [17] Heisser, T., Cardoso, R., Guo, F., Moellers, T., Hoffmeister, M., & Brenner, H. (2021). Strongly divergent impact of adherence patterns on efficacy of colorectal cancer screening: the need to refine adherence statistics. Clinical and Translational Gastroenterology, 12(9). DOI: <u>https://doi.org/10.14309%2Fctg.000000000000399</u>
- [18] Kumar, R., Smith, K. S., Deng, M., Terhune, C., Robinson, G. W., Orr, B. A., ... & Northcott, P. A. (2021). Clinical outcomes and patient-matched molecular composition of relapsed medulloblastoma. Journal of Clinical Oncology, 39(7), 807. DOI: <u>https://doi.org/10.1200%2FJCO.20.01359</u>
- [19] Holtgrave, D. R., Barranco, M. A., Tesoriero, J. M., Blog, D. S., & Rosenberg, E. S. (2020). Assessing racial and ethnic disparities using a COVID-19 outcomes continuum for New York State. Annals of epidemiology, 48, 9-14. DOI: <u>https://doi.org/10.1016/j.annepidem.2020.06.010</u>
- [20] He, Y., Wang, J., Li, F., & Shi, Y. (2020). Main clinical features of COVID-19 and potential prognostic and therapeutic value of the microbiota in SARS-CoV-2 infections. Frontiers in Microbiology, 11, 1302. DOI: <u>10.1016/j.immuni.2012.04.011</u>
- [21] Koyuncu, I., Kocyigit, A., Ozer, A., Selek, S., Kirmit, A., & Karsen, H. (2018). Diagnostic potential of Brucella melitensis Rev1 native Omp28 precursor in human brucellosis. Central European Journal of Immunology, 43(1), 81-89. DOI: <u>https://doi.org/10.5114/ceji.2018.74877</u>