Seroprevalence of bovine brucellosis and associated risk factors among smallholder dairy cattle farmers in Hai and Meru District Councils Northern Tanzania

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Abstract
Brucellosis is an endemic zoonotic disease caused by a facultative intracellular gram-negative bacterium of the genus Brucella. This study was carried out to establish the current status of brucellosis and possible associated risk factors in the smallholder dairy cattle in Hai and Meru District. A cross-sectional study was conducted between January and June 2022 to investigate the current seroprevalence of bovine brucellosis and possible associated risk factors in small dairy farms in the study areas. A total of 400 cattle was sampled with blood collected from 10 villages in each district. Blood samples were analyzed for Brucella circulating antibodies using the Rose Bengal Plate Test and c-ELISA. A structured questionnaire was administered to 200 smallholder dairy cattle farmers to determine the potential risk factors associated with brucellosis among dairy cattle in the study areas. The overall seroprevalence of bovine brucellosis in the study area was 0.50% and 0% for the Hai and Meru districts, respectively. Analysis of knowledge and management practices of brucellosis in the study area showed that the majority of farmers 74.5%, (149/200) were knowledgeable about the name of the disease; though, the majority 87.9%, (131/149) did not know the clinical signs. Most likely, awareness and biosecurity based on the nature of the farming system (zero grazing) contributed to the low seroprevalence; thus, none of the risk factors were associated with the disease. Therefore, under the smallholder dairy farming system, a four-stage roadmap for progressive control of brucellosis in animals and humans as recommended by FAO could be implemented with farming system modifications to eradicate the disease.

Keywords: Brucellosis, dairy cattle; seroprevalence; risk factors; Tanzania

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Introduction

The smallholder dairy sector is among the growing livestock subsectors in Tanzania. This sector plays a great role in the national food security, with its importance ranging from nutrition, health, employment opportunity; manure for farm fertilization, source of energy through biogas production; and generation of household income through selling of milk surplus and milk products (URT 2015). Nevertheless, the dairy sector in Tanzania is constrained by a number of challenges, such as a seasonal change in pasture availability and quality, lack of broad-based dairy production technologies, low rate of milk processing, poor milk quality, poor milk handling facilities, long calving interval and diseases including zoonoses such as anthrax and brucellosis (Bingi & Tondel., 2015, Maleko et al., 2018).

Brucellosis is among the endemic-neglected zoonotic diseases of social-economic importance in many regions and countries, such as Central Asia, India, Near East countries, Mexico, South and Central America, European Mediterranean countries and African countries, including Tanzania (Corbel 2006). Globally, the prevalence of bovine brucellosis ranges between 2% and 36% (Seleem et al., 2010; Abu Sulayman et al., 2020; Bodenham, 2020; Khurana et al., 2021; Holt et al., 2021; Ntivuguruzwa et al., 2020; Wainaina et al., 2020; Djangwani et al., 2021; Mengele et al., 2023). However, the disease has been eradicated in some European countries because of successful eradication and control programmes, including active surveillance coupled with test and slaughter policy and mass vaccination of domesticated animals (Seleem, Boyle, and Sriranganathan 2010). In Tanzania, the disease is among the six prioritized zoonotic diseases that require national attention. The endemicity of bovine brucellosis in Tanzanian dairy farming is attributed to a number of risk factors, such as inadequate surveillance of the disease, lack of vaccination of programmes, unrestricted animal replacement and movements, weak regulatory framework in the culling of brucellosis-positive reactors, poor knowledge of the disease, improper disposal of aborted fetus and retained placentas, animal interactions in grazing and watering points and breeding practices (Asakura et al., 2019; Sagamiko, 2019; Bodenham et al., 2020; Ntirandekura et al., 2021; Katandukila et al., 2021).

Brucellosis infection in smallholder dairy cattle causes not only negative social-economic impacts but also public health consequences (Akakpo 2009). Some efforts to prevent and control the spread of brucellosis in Tanzania under smallholder dairy farming system were effectively practiced early in 1980-1990 when prevalence was reduced to ≤ 2% in dairy cattle. However, recent studies have showed that the seroprevalence of brucellosis in smallholder dairy cattle is re-emerging from 0%-22.1% in various hotspot areas of Tanzania following the collapse of Tuberculosis and Brucellosis control Programme in animals (Swai E, Nkya R and Kambarage, 2000; Karimuribo et al., 2007; Mellau & Wambura, 2009; Shirima et al., 2010; Swai & Schoonman, 2010; Shirima et al., 2016; Bodenham, 2020; Mengele et al., 2023). Despite the smallholder dairy in the northern Tanzania supplying milk to major milk processing plants; little work has been done to ascertain the current status of brucellosis in dairy cattle in this area. Therefore, this calls for a detailed investigation to establish the
disease status, spread, hot spots and possible associated risk factors in smallholder dairy cattle in the Hai and Meru District Councils representing the Kilimanjaro and Arusha regions where smallholder dairy farming is highly prominent.

Materials and Methods

Study area

The research work was carried out in the Hai and Meru District Councils in northern Tanzania (Fig.1). The two districts were selected based on the high number of smallholder dairy cattle farms. Hai has a total of 49,316 dairy cattle and occupies an area of 1,011 km$^2$ (101,100 ha), while Meru has a total of 98,001 dairy cattle and an area of 1,268.2 km$^2$.

Figure 1. Map of the study area showing the boundaries of Meru and Hai Districts.
Source: Created by using Quantum Geographic Information System (QGIS).

Study design, sample size and sampling procedures

A cross-sectional study was carried out between January and June 2022 involving adult dairy cattle of at least one year of age under the smallholder farming system. The sample size was calculated using Fisher’s (1998) statistical procedure, with a confidence interval of 95%, a margin error of 5% and an estimated prevalence of 50% in smallholder dairy cattle.

The following formula was used for the sample size calculation:

$$ N = \frac{Z^2 \times P (1 - P)}{C^2} \quad (eq \ 1) $$

where $N$=Sample size, $P$=Estimated prevalence = 0.5, $Z$=Level of confidence as 1.96 and $c$ = Desired precision level = 0.05.

The calculated minimum sample size was 384. However, the sample size “$N$” used in this study was 400 smallholder dairy cattle. The study involved both purposive and multistage random sampling procedures. These sampling procedures involved 62 villages and 17 wards in the Hai District Council, while there are 90 villages and 26 wards in the Meru District Council. The study involved 10 villages and 5 wards from each District. The Wards inclusion criteria in this study was possession of at least 500 dairy cattle. First, there was a purposive selection of 5 wards with a large number of dairy cattle from each district, followed by a multistage random sampling of two villages from each ward in which 20 dairy cattle from 10 dairy cattle-keeping
households in each Village (2 dairy cattle @ household) were selected randomly for blood collection. At every stage of random sampling, run if () function in statistical software R was deployed.

**Field data collection**

Cattle were restrained manually and humanly to avoid harm or any causes of animal discomfort during sample collection. The exercise was performed in compliance with the Tanzania Animal Welfare Act, part V (Animal Welfare Act 2008). Using a halter, the animal’s head was fastened to an elevated position to allow visibility of the jugular vein. Then, the thumb finger was pressed at the base of the jugular groove to raise blood pressure and visualize the vein by blocking the vein. With a plain vacutainer and needle, 10 ml of blood was drawn from the jugular vein. Each animal was identified according to the identity type provided by the owner for subsequent identification. This enables proper labelling of the vacutainer tubes containing blood samples. To avoid albumin coagulation, which prevents serum formation during the centrifugation process, the blood samples were left at ambient temperature for approximately 30 minutes before centrifugation. Centrifugation was performed at the Tanzania Veterinary Laboratory Agency (TVLA) in the northern zone office, Arusha, where the vacutainer tubes were spun at 3000 rpm for 10 minutes (BHG S Segurita-Germany). After centrifugation, the tubes were removed, and the sera were decanted into 2.0 ml cryogenic vials. The sera were stored temporarily at the TVLA laboratory at -20°C soon after separation before transfer to The Nelson Mandela African Institution of Science and Technology’s (NM-AIST) laboratory for analyses. Likewise, at the NM-AST laboratory, the sera were stored at -20°C before execution of analysis.

**Questionnaire survey**

To investigate the possible risk factors for brucellosis in smallholder dairy cattle, a paperless questionnaire survey comprising both closed- and open-ended questions was developed and coded using the Open Data Kit [https://opendatatoolkit.org/(ODK)] mobile application by the investigator. The questionnaire covered a wide range of information, such as abortion cases, recent reproductive status, history of retained placenta, knowledge about bovine brucellosis, use of vaccines, use of veterinary services, herd management practices, herd size, handling of aborting cows and aborted fetus, heifer sources, breeding methods, grazing system, milk production trends, past two calving dates, livestock movement and interaction with neighboring cattle herds as well as milk distribution channels, price and value chain (Appendix 1). Pretesting of the questionnaire was performed in smallholder dairy cattle-keeping households in Monduli District before the development of the final version. During the field visit, the questionnaire was administered to the respondents (head of household or someone knowledgeable with the herd and above 18 years old). It took approximately 25-30 minutes to complete the questionnaire successfully.

**Serological analysis**

Laboratory analysis was carried out at The Nelson Mandela African Institution of Science and Technology laboratory, where a Rose Bengal Plate Test (RBPT) was used as a screening test for all 400 serum samples, while the competitive enzyme-linked immunosorbent assay (cELISA) was used for confirmation of the positive reactors.
Through RBPT, all 400 serum samples were screened for Brucella antibodies using the Zoetis \textsuperscript{TM} (Rev 4/2017) Brucella Rose Bengal test kit from Delpharm Biotech (Lyon Cedex 07-France). Briefly, according to the manufacturer’s instructions, equal volumes (30 μL) of the test serum and antigen were mixed thoroughly on the glass plate using an applicator stick, and the plate was gently hand rocked to allow mixing. After four minutes, the plates were visually examined for agglutination in comparison with a positive control. Any degree of agglutination was considered positive, while the absence of agglutinates was considered negative. The results were recorded and saved in a Microsoft Excel spreadsheet. The white glass plate was washed with clean water and methylated spirit and then dried for approximately 5-10 minutes before reuse.

The serum sample that reacted positive to RBPT was tested by cELISA according to (COMPELISA Rev01/2020) APHA SCIENTIFIC (Animal and plant Health Agency); New Haw Addlestone, Surrey, KT15 3NB United Kingdom. The only one RBPT positive reactor and 39 serum samples that reacted negatively on RBPT were randomly selected to make a single full microtitre plate during the cELISA test.

Data storage and statistical analysis

The obtained data were saved in Microsoft Excel spreadsheet software which was also used for descriptive analysis. The rest of data statistical analysis were done by using R statistical software version 4.0.3 (2020-10-10). The prevalence of brucellosis was determined using descriptive analysis based on the RBPT and cELISA tests using the following equation.

\[ p_i = \frac{x_i}{n_i} \] (eq 2)

where \( x_i \) is the number of animals testing positive for brucellosis in a given administrative area, and \( n_i \) is the total number of animals tested in that administrative area. Furthermore, 95% exact binomial confidence intervals (CIs) were calculated using the binom.test function from the core R (www.R-project.org) stats package. The same formula was used to compute the overall seroprevalence across the study area.

In univariate analysis, the chi-square test was used to compare two or more proportions to determine the degree of relationship and significant differences. The variables that had \( P < 0.05 \) at the 95% confidence interval in univariable analysis were included in multivariate analysis, where the odds ratio (OR) was utilized to investigate the relationship between brucellosis risk variables and brucellosis prognosis.

**Ethical Clearance**

The research project was approved by Kibong’oto Infectious Disease Hospital-The Nelson Mandela African Institution of Science and Technology and the Centre for Education Development in Health, Arusha (KIDH-NM-AIST-CEDHA)-KNCHREC) ethical committee with certificate number KNCHREC0067/04/2022 issued on 27th June 2022. Additionally, permission was granted by the respective District Executive Directors (DEDs) in response to the introduction letters from NM-AIST. The willingness of the head of households to participate in this study was sought through written consent before the execution of the study.

**Results**

**The sociodemographic profile of the respondents**

The sociodemographic profile of the respondents indicated that 72%, (144/200)
were males and 28%, (56/200) were females of which the majority of the respondents (81%; 162/200) were aged between 41 and 60 years (Table 1).

<table>
<thead>
<tr>
<th>Variables</th>
<th>Category</th>
<th>Meru</th>
<th>Hai</th>
<th>Total (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gender</td>
<td>male</td>
<td>67</td>
<td>77</td>
<td>144 (72)</td>
</tr>
<tr>
<td></td>
<td>Female</td>
<td>33</td>
<td>23</td>
<td>56 (28)</td>
</tr>
<tr>
<td>Age</td>
<td>18-25</td>
<td>1</td>
<td>0</td>
<td>1 (0.5)</td>
</tr>
<tr>
<td></td>
<td>26-40</td>
<td>6</td>
<td>8</td>
<td>14 (7)</td>
</tr>
<tr>
<td></td>
<td>41-60</td>
<td>79</td>
<td>83</td>
<td>162 (81)</td>
</tr>
<tr>
<td></td>
<td>&gt;60</td>
<td>13</td>
<td>11</td>
<td>24 (12)</td>
</tr>
<tr>
<td>Marital status</td>
<td>Single</td>
<td>11</td>
<td>9</td>
<td>20 (10)</td>
</tr>
<tr>
<td></td>
<td>Married</td>
<td>88</td>
<td>83</td>
<td>171 (85.5)</td>
</tr>
<tr>
<td></td>
<td>Divorced</td>
<td>1</td>
<td>0</td>
<td>1 (0.5)</td>
</tr>
<tr>
<td></td>
<td>Widowed</td>
<td>3</td>
<td>5</td>
<td>8 (4)</td>
</tr>
</tbody>
</table>

Respondents’ knowledge and awareness of brucellosis
Although 74.5% (149/200) of the respondents were aware of the disease (brucellosis) majority (87.9%; 131/149) were ill of the disease clinical signs in animals. Among the clinical signs of the disease, abortion was mentioned by 12.08% (18/149) of the respondents, while no one knew the rest of the clinical signs of brucellosis (Fig. 2).

![Figure 2. Awareness of the respondents on the clinical signs of cattle brucellosis in the study areas.](image-url)
The results also showed that none of the respondents heard of brucellosis vaccine (S19) has never used in the study area whereas, anthrax vaccine was well known by majority of respondents (Fig.3)

![Figure 3. Respondents’ awareness on commonly used vaccines in dairy cattle](image)

**Disposal methods of the aborted fetus and retained placenta.**

The results show that 7.5%, (15/200) of respondents’ herd had a history of abortion while 1.5% (3/200) reported retained placenta cases. Majority of the respondents (80%, 12/15) buried aborted foetus and retained placenta while 20%, (3/15) thrown to dogs (Fig.4)

![Figure 4. Disposal methods practiced by respondents on the aborted fetus and retained placenta](image)

**Animal breeding methods used in the study area.**

Majority of the respondents (55%, 110/200) use artificial insemination compared to 32%, (64/200) who use bulls for breeding (Fig. 5), Of these who are using bulls, 64%, (41/64) hire from another smallholder dairy farms. However, opinions on choice decision were based on cost (51.5%, 103/200), accessibility (34%, 68/200) and efficiency of the service type (14.5%, 29/200).
Grazing system and animal interactions. The majority of respondents (87.5%, 175/200) practise zero grazing system with only 1%, (2/200) practised a free-range system (Fig. 6 Furthermore, 99%, (198/200) of the respondents reported that their cattle were neither herded nor fed together with sheep and goats.

Animal sources for heifer replacement. The results show that more than sixty percent of the respondents (129/200) acquired heifers from other smallholder dairy cattle farms with the district for replacement, whereas 35% (69/200) used to upgrade female calves to become heifers from their own farm. (Fig. 7)
Management of animals that have failed to conceive.

Although majority of farmers (90.5%, 181/200) reported high conception rate by using both artificial insemination and bull service, few farmers with animals that failed to conceive sold to other smallholder farmers as reported by 63.16%, (12/19) of the respondents (Fig. 8).

Seroprevalence of bovine brucellosis in smallholder dairy cattle in the Hai and Meru districts

A total of 400 dairy cattle from smallholder farms were tested for brucellosis using RBPT and cELISA tests. The animal-level seroprevalence of brucellosis in Meru and Hai Districts was 0% and 0.5%, respectively, while the herd-level seroprevalence was 0% and 1% in Meru and Hai Districts, respectively. However, the difference between the two
districts was not statistically significant at p<0.05 and 95% CI. The overall animal- and herd-level seroprevalence of brucellosis in the study area was 0.25% and 0.5%, respectively. (Table 2).

Table 2. The individual and overall seroprevalence (%) of bovine brucellosis in the Hai and Meru District Councils

<table>
<thead>
<tr>
<th>Study District</th>
<th>Animal level seroprevalence</th>
<th>Herd level seroprevalence</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>N (n)</td>
<td>Prevalence (%)</td>
</tr>
<tr>
<td>Meru</td>
<td>200 (0)</td>
<td>0</td>
</tr>
<tr>
<td>Hai</td>
<td>200 (1)</td>
<td>0.5</td>
</tr>
<tr>
<td>Overall results</td>
<td>400 (1)</td>
<td>0.25</td>
</tr>
</tbody>
</table>

N = Number of animals tested  n = Number of positive reactors

Association of the seroprevalence of bovine brucellosis at animal level risk factors.

Analysis of animal level risk factors and seroprevalence of bovine brucellosis among smallholder dairy cattle showed that the animal that tested positive was an adult Friesian female with neither abortion history nor infertility problems as well as history of retained placenta. Also, the positive reactor animal had no history of retained placenta. All animal levels risk factors (Age, sex, breed, abortion history, retain placenta and failure to conceive) were statistically insignificant (p values > 0.05) under univariable analysis (Table 3).

Table 3. Association of the seroprevalence of bovine brucellosis at animal level risk factors

<table>
<thead>
<tr>
<th>Risk factors</th>
<th>Category</th>
<th>N (n)</th>
<th>Prevalence (%)</th>
<th>Univariable analysis</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age</td>
<td>Young</td>
<td>22(0)</td>
<td>0</td>
<td>1.51e-28</td>
</tr>
<tr>
<td></td>
<td>Adult</td>
<td>378(1)</td>
<td>0.25</td>
<td></td>
</tr>
<tr>
<td>Sex</td>
<td>Male</td>
<td>23(0)</td>
<td>0</td>
<td>9.95e-32</td>
</tr>
<tr>
<td></td>
<td>Female</td>
<td>377(1)</td>
<td>0.25</td>
<td></td>
</tr>
<tr>
<td>Breed</td>
<td>Jersey</td>
<td>33(0)</td>
<td>0</td>
<td>0.30369</td>
</tr>
<tr>
<td></td>
<td>Ayrshire</td>
<td>61(0)</td>
<td>0</td>
<td>0.8591</td>
</tr>
<tr>
<td></td>
<td>Friesian</td>
<td>306(1)</td>
<td>0.25</td>
<td></td>
</tr>
<tr>
<td>History of abortion</td>
<td>Yes</td>
<td>29(0)</td>
<td>0</td>
<td>1.72e-29</td>
</tr>
<tr>
<td></td>
<td>No</td>
<td>371(1)</td>
<td>0.25</td>
<td></td>
</tr>
</tbody>
</table>
### Table 4. Association of the seroprevalence of bovine brucellosis at herd level risk factors

<table>
<thead>
<tr>
<th>Risk factor</th>
<th>Category</th>
<th>N (n)</th>
<th>Prevalence (%)</th>
<th>(\chi^2)</th>
<th>p value</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Grazing system</strong></td>
<td>Free range</td>
<td>4(0)</td>
<td>0</td>
<td>0.14322</td>
<td>0.9309</td>
</tr>
<tr>
<td></td>
<td>mixed</td>
<td>46(0)</td>
<td>0</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>zero grazing</td>
<td>350(1)</td>
<td>0.25</td>
<td></td>
<td></td>
</tr>
<tr>
<td><strong>Type of service</strong></td>
<td>AI</td>
<td>224(0)</td>
<td>0</td>
<td>2.155</td>
<td>0.3404</td>
</tr>
<tr>
<td></td>
<td>Bull</td>
<td>126(0)</td>
<td>0</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Both</td>
<td>50(1)</td>
<td>0.25</td>
<td></td>
<td></td>
</tr>
<tr>
<td><strong>Heifer source</strong></td>
<td>livestock markets</td>
<td>2(0)</td>
<td>0</td>
<td>0.01861</td>
<td>0.8915</td>
</tr>
<tr>
<td></td>
<td>smallholder dairy cattle farms</td>
<td>258(0)</td>
<td>0</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Big dairy cattle farms</td>
<td>2(0)</td>
<td>0</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>own farm</td>
<td>138(1)</td>
<td>0.25</td>
<td></td>
<td></td>
</tr>
<tr>
<td><strong>Frequent contacts with other herds</strong></td>
<td>Yes</td>
<td>2(0)</td>
<td>0</td>
<td>8.24e-23</td>
<td>1</td>
</tr>
<tr>
<td></td>
<td>No</td>
<td>398(1)</td>
<td>0.25</td>
<td>0</td>
<td>1</td>
</tr>
<tr>
<td><strong>Disease name</strong></td>
<td>Yes</td>
<td>299(1)</td>
<td>0.25</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>No</td>
<td>101(0)</td>
<td>0</td>
<td></td>
<td></td>
</tr>
<tr>
<td><strong>Brucellosis diagnosis</strong></td>
<td>Yes</td>
<td>11(0)</td>
<td>0</td>
<td>2.56e-26</td>
<td>1</td>
</tr>
<tr>
<td></td>
<td>No</td>
<td>389(1)</td>
<td>0.25</td>
<td></td>
<td></td>
</tr>
<tr>
<td><strong>Vaccinated against brucellosis</strong></td>
<td>Yes</td>
<td>0(0)</td>
<td>0</td>
<td>8.38e-27</td>
<td>1</td>
</tr>
<tr>
<td></td>
<td>No</td>
<td>400(1)</td>
<td>0.25</td>
<td></td>
<td></td>
</tr>
<tr>
<td><strong>Access to veterinary services</strong></td>
<td>Yes</td>
<td>(0)</td>
<td>0</td>
<td>1.26e-26</td>
<td>1</td>
</tr>
<tr>
<td></td>
<td>No</td>
<td>44(1)</td>
<td>0.25</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

N = Number of animals tested; n = Number of positive reactors; RP = Retained placenta

**Detailed information of the seropositive cattle**

Further information revealed that the seropositive cattle was a Friesian adult female kept under zero grazing system. The herd had 4 dairy animals with no small ruminants. Natural mating and artificial insemination were interchangeably used to service the animals in the herd. The knowledge and practises related to bovine brucellosis and husbandry with the herd owner did not differ from the rest of farmers. All herd level variables (grazing system, type of service, source of heifers, contacts and vaccination) were not associated with the brucellosis seropositivity (p values >0.05 (Table 4).
Discussion

Brucellosis is among the six prioritized zoonotic disease in Tanzania that has social economic effects. This study was carried to establish the current status of brucellosis and possible associated risk factors in the smallholder dairy cattle in Hai and Meru District.

Seroprevalence of bovine brucellosis among smallholder dairy cattle.

The findings from this study, indicate that the sampled cattle in Meru district did not show antibodies against brucella infection, however, one dairy cattle in Hai district indicated to have been exposed to brucella infection. This indicates that the two districts are not among the brucellosis hotspots as reported elsewhere (Mengele et al., 2023; Shirima et al., 2018). Similarly, in this study, it was found that dairy farmers are not using the S19 vaccine against brucellosis and this indicates that the infected animal was due to natural exposure of the pathogen. The low seroprevalence of bovine brucellosis in the study area lies within the range of brucellosis studies in dairy cattle carried out in several parts of Tanzania (Mengele et al., 2023; Mdegela et al., 2004; Karimuribo et al., 2007; Alexander, 2017; Mhozya, 2017). Similarly, the findings of this study are in agreement with other studies conducted across the World that indicate a low seroprevalence in smallholder dairy cattle (Kothowa et al., 2021, (Hesterberg et al. 2008), (Getahun 2021), (Nguna et al. 2019), (Hassan et al. 2014). The results of this study are contrary to the findings of other studies where the seroprevalence was very high (Swai & Schoonman, 2010 ; Mengele et al., 2023). The difference in seroprevalence in various studies in Tanzania can be caused by variations in study design, farming systems, management practices and other biosecurity measures taken by farmers. The very low seroprevalence obtained in this study might be attributed by a number of factors such as zero grazing system, animal replacement practices, disposal method of the aborted foetus and breeding methods.

In this study, it was found that zero grazing system was the dominant grazing system practiced by most of the smallholder dairy cattle farmers. Through zero grazing system, animals are fed on fodder using cut and carry practices. Therefore, there is less interactions of animals between herds thus acts as one of the key biosecurity control options against disease transmission including brucellosis. This observation agrees with the findings of other studies (Karimuribo, 2007 ; Swai & Schoonman, 2010) that zero grazing system minimize the level of infection since animals from different herds do not interact to each other. This is in contrary to other farming systems such as pastoral and agro-pastoral systems where herds with multispecies interact frequently in grazing and watering points thus perpetuates disease transmission (Assenga et al., 2015; Shirima, 2005). Furthermore, the zero grazing system practised in the study area separates small ruminants from being herded together with the dairy cattle thus minimize the cross-infection risk as well (Shirima, 2005 ; Rubegwa, 2015 ; Oromia et al., 2022 ; Mengele et al., 2023).
Based on the fact that both districts practised dairy farming for decades, animal replacement becomes feasible from within minimising introduction of animals from outside herds/districts. another reason for the low seroprevalence observed in this study. It was found that most animals for replacement were originated from either within the herds or between the herds in the study area. The mode of animal acquisition in the study area does not favour the introduction of animals that might be infected from other areas. This observation concurs with the findings from other studies (Shirima, 2005; Karimuribo, 2007; Alexander, 2017). However, the findings of this study are not similar to that of Rubegwa et al., (2015) and Adera et al., (2019) who reported higher bovine seroprevalence in homebred animals than brought animals.

In this study, it was found that most common breeding method used was artificial insemination instead of sharing bulls. This type of breeding method could contribute to the observed low seroprevalence of brucellosis in the sampled cattle. The disease’s free semen used in artificial insemination limits possible transmission of Brucella pathogens from the infected bulls to cows. This findings from the study areas is also supported by others elsewhere (Corbel, 2006; Shirima, 2005) and Mfune, 2015) who reported that there is high prevalence of Brucella in areas with low rate of using artificial insemination breeding method. Although proportion of dairy cattle farmers used bulls for mating; they are sourced from within the district and when screened were negative to both RBPT and cELISA. The use of bulls is not uncommon in pastoral farming systems thus may attribute to the level of infection reported compared to dairy cattle farming (Mellau and Wambura, 2009; Swai and Schoonman, 2010; Nguna et al., 2019).

Reports of abortion incidences and retain placentas in the current study may be clear evidence of low reproductive diseases including brucellosis. From this study, it was found that abortion and retained placenta were not common cases to happen in dairy cattle. However, cases of aborted foetus and retained placenta were disposed of properly by burying them onto the ground. This further prevents disease perpetuation as reported earlier that feeding raw to dogs amplify spread of the disease (Shirima 2005; Sijapenda et al., 2017; Ntirandekura et al., 2018; Ismail et al., 2019 Ntivuguruzwa et al., 2020; Mengele et al., 2023).

**Animal and herd level factors associated with brucellosis in smallholder dairy cattle**

Both animal level (sex, age, breed type, history of retained placenta and abortion) and herd level (farmer’s awareness of the disease name, clinical signs of brucellosis in cattle, proper disposal methods of the aborted fetus, management of infertile animals, grazing system, breeding method, animal replacement, animal housing system, animal vaccination) risk factors were not significantly associated with the seropositivity due to a very low seroprevalence of the sampled cattle in this study. Since this happened in the absence of any formal prevention and control interventions it calls for a systematic surveillance and monitoring to maintain the status. Farms brucellosis certification coupled with continues education and awareness may be a strategic approach.

**Conclusion and recommendations**

In this study, it was found that the sampled cattle had a very low
seroprevalence. Likely, brucellosis prevalence in smallholder dairy cattle in Hai and Meru districts is relatively low. Zero grazing system coupled with in-house breeding and animal replacement from within the study areas may have provided biosecurity measures for brucellosis spread thus, need to be intensified to limit introduction of infections in the study areas.

Based on the findings from this study; it is recommended that; the surveillance monitoring approach may shift to bulk milk sampling to detect exposed herds in the event of low seroprevalence. Also, Knowledge about brucellosis is highly recommended as it can guide control efforts and improve information about local risk factors as well as the extent of dairy farmers’ understanding of the disease to foster creation of better extension campaigns on the brucellosis. Importantly, brucellosis certification scheme initiated after this study be strengthened and monitored to ensure a disease-free area.

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