

Isolation, Identification and Antimicrobial Resistance Profile of Staphylococcus in Dairy Cows of Assosa town, Abrahamo, Ura and Bambasi Districts of Benishangul Gumuz Regional State, Western Ethiopia

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ABSTRACT: A cross - sectional study was conducted from November 2023 to July 2024 in Dairy cattle in Assosa town, Abrahamo, Ura and Bambasi districts in order to estimate the prevalence of mastitis, isolate and identify *S.aureus* from Dairy cows, assess its antimicrobial resistance pattern and identify risk factors associated with mastitis. A total of 385 Dairy cows milk samples were collected with random sampling techniques. In this study, out of the total lactating cows examined, 150(38.96%) mastitis prevalence was found to be affected with mastitic infection. During laboratory examination, 85/385(22.07%) of the *S.aureus* was isolated and 69/385(17.92%) of other coagulase negative *staphylococcus spp (CNS)* were identified. The relative proportional prevalence of *Staphylococcus aureus* was 85/150(56.66%). They were found to be statistically significant ($P<0.00$). The highest mastitic dairy cows' distribution were observed in Abrahamo (51.9%) while the lowest prevalence was seen in Ura (29.33%). In this study, breed, age, parity, tick infestation, teat lesion were non- significant ($P>0.05$) while length of lactation, previous mastitis history, blind teat, previous mastitis treatment history, milk hygiene and floor type were significant ($p<0.05$). The present result showed a significant association of resistance pattern with *S.aureus* isolates, particularly to penicillin G (78.84%), Cefoxitin (76.92%), Tetracycline (69.23%), Streptomycin (61.53%) and Gentamycin (53.84%) were investigated. Hence, regular resistance follow-up, using antimicrobials sensitivity tests helps to select effective antibiotics and to reduce the problems of drug resistance developments towards commonly used antimicrobials so as to reduce the problem encountered.

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1. INTRODUCTION

Ethiopia has the largest cattle population in Africa with an estimated population of 52.13 million (CSA, 2007) and contributes 40% to the annual agricultural output, and 15% total gross domestic product. Cattle produce a total of 1.5 million tonnes of milk and 0.331 million tonnes of meat annually (FAO, 2005). Cows represent the biggest portion of cattle population of the country, around 42% of the total cattle heads are milking cows (CSA, 2008). However, milk production often does not satisfy the country's requirements due to a multitude of factors. Mastitis is among the various factors contributing to reduced milk production (Biffa *et al.*, 2005).

Staphylococcosis, an infectious bacterial zoonosis of global significance, is caused by *S. aureus*, which is gram- positive, non-capsulated, non-motile, catalase positive, non-sporulated organism, grape-like clusters, 0.5-1.5 micrometer in diameter (Harris *et al.*, 2002). Pathogenic *Staphylococci* are commonly identified by their ability to produce coagulase, and thus clot blood. This distinguishes the coagulase positive strains, *S. aureus*, *S. intermedius* and *S. hyicus* from the other Staphylococcal species such as *S.*

epidermidis that are coagulase-negative (Harris *et al.*, 2002). *S. aureus* is both commensal and pathogen. It is found as a commensal associated with skin, skin glands and mucous membranes. *S. aureus* affects skin, soft tissues, bloodstream and lower respiratory tract. It also causes severe deep-seated infections like endocarditis and osteomyelitis (Schito, 2006). *S. aureus* also causes severe animal diseases, such as suppurative disease, arthritis and urinary tract infections (Lowy, 1998).

S. aureus is present in a variety of locations in the dairy farms, in many occasions it was isolated from swabs taken from the cows head, skin swabs, legs and nasal mucosa (Zadoks *et al.*, 2000). Furthermore *S. aureus* was found on the milkers' hands as well as on the nasal mucous membrane of the humans working at the dairy farms, in bedding and the drinkers (Benić *et al.*, 2012). However an infected udder quarter remains the main reservoir of the bacteria, which transmitted mostly during the milking time. Recent researches show that many biotypes and genotypes exist on the dairy farms (Zadoks *et al.*, 2002; Smith *et al.*, 2005).

S. aureus plays its most significant animal pathogenic role as cause of intramammary infections in cattle and small ruminants leading to considerable economic losses in dairy farms. The pathogen is frequent causative agent of clinical or subclinical mastitis in cattle (Asperger and Zangerl, 2003). Presence of *S. aureus* on the skin and mucosae of food producing animals, such as ruminants, and the frequent association of the pathogen with mastitis, often leads to contamination of milk (Jablonski and Bohach, 1997). Contamination of milk can also occur from environmental sources during handling and processing (Peles *et al.*, 2007). Milk is a good substrate for *S. aureus* growth and dairy products are common sources of staphylococcal food-poisoning (Morandi *et al.*, 2007).

Enterotoxin-producing *S. aureus* plays an important role as causative organism of food intoxications. In many countries, *S. aureus* is considered to be the second or third most common pathogen causing outbreaks of food poisoning only outnumbered by *Salmonella* species, and in competition with *Clostridium perfringens* (Aycicnek *et al.*, 2001).

Although a variety of antibiotics can be used against this organism, *S. aureus* mastitis has been found to respond poorly to antibiotic treatment (Barkema *et al.*, 2006). The increased resistance of *S. aureus* isolates to several antimicrobial agents has been reported (Gentilini *et al.*, 2000). The determination of antimicrobial susceptibility of clinical isolates is required not only for therapy but also for monitoring the spread of resistant strains throughout the populations. β -lactam antibiotics are the most frequently used in intramammary infusion therapy. Bacterial resistance mechanisms to this class of antibiotics include production of β -lactamase and low-affinity penicillin-binding protein 2a (PBP 2a) determined by the presence of the chromosomal gene *mecA*. The latter, designated for methicillin resistance, precludes therapy with any of the currently available β -lactam antibiotics, and may predict resistance to several classes of antibiotics (Moon *et al.*, 2007).

The usage of antibiotics correlates with the emergence and maintenance of antibiotic resistant traits within pathogenic strains (Shitandi and Sternesjo, 2004). These traits are coded for by particular genes that may be carried on the bacterial chromosome, plasmids, and transposons or on gene cassettes that are incorporated into integrons (Rychlik, 2006), thus are easily transferred among isolates. Multiple antibiotic resistant *S. aureus* strains have been isolated from milk obtained from cattle samples in many parts of the world (Pesavento *et al.*, 2007).

From a number of epidemiological studies of Staphylococcal mastitis conducted, only few of them were done on economic and zoonotic significance *S. aureus* from milk samples in Benishagul Gumuz region. In Benishagul Gumuz region, there are few studies in and Around Asossa town and also Bambasi District (Asmamaw A *et al.*, 2017, 2018). And hence, knowledge of zoonotic and economic impact of *S. aureus* and treatment failure in developing countries is necessary to make decisions and prerequisite for establishing control strategies.

So far, there was no study done on to assess the epidemiology of *Staphylococcus spp* in Assosa town, Abrahamo, Ura and Bambasi districts.

Therefore, the objectives of the current study are:

- To determine the prevalence of bovine mastitic *S. aureus*
- To isolate and characterize *S. aureus* from mastitic lactating cows
- Assessment of the risk factors associated with *Staphylococcus* infections
- To determine the antimicrobial resistance pattern of *S. aureus* species

2. MATERIALS AND METHODS

2.1 Study Area

The study were conducted in Assosa town, Abrahamo, Ura and Bambasi districts. Assosa is the capital city of the Benishangul-Gumuz Regional State and composed of 74 administrative peasant associations, which is located at 8°30' and 40°27' N latitude and 34°21' and 39°1' E longitude 687 kms Northwest of Addis Ababa (CSA, 2015). The altitude of Asossa ranges from 580 to over 1544 meter above sea level. The area is characterized by low land plane agro- ecology according to National Meteorological Service Agency (NMSA, 2014) with average annual rainfall of 1316 mm with uni-modal type of rainfall that occurs between April and October. Its mean annual temperature ranges between 16.75°C and 27.9°C. Asossa zone has 35.6% of the livestock population of the region constituting 61, 234 cattle, 191, 83 goats, 19,729 sheep, 25,137 donkeys, 439,969 poultry and 73,495 beehives (CSA, 2015), and the Assosa District has 16,990 cattle, 30,728 shoat, 57,089 poultry and 5,240 donkey (Bureau of agriculture, 2016).

Bambasi district has 38 kebeles stretches over an area of 2210.16 square k.m with human population of 62693. The region is found in the north west of the country between latitude of 9 and 11°N and longitude of 34 and 35°E and its altitude range is 1500-1900 meter above sea level. Annual rain fall is between 1350-1400 mm with uni modal type of rain

Asossa, for microbiological examination. If immediate inoculation is not convenient, samples were kept at 4°C until cultured for isolation.

2.6. Study Methodology

2.6.1. Questionnaire survey

Data on each sampled cow was collected using a properly designed questionnaire format for determining the associated risk factors. This includes milker status, environmental contamination, age, body condition, parity, and stage of lactation, breed, previous history of mastitis treatment, barn floor type, milking hygiene, milking practice and other relevant information related to other management practices related to mastitis will be gathered. Udder and milk abnormality (injuries, swelling, milk clots and abnormal secretions, etc) were also recorded. Drug usage practice in the study area will be also collected to evaluate its contribution to the emergence of antimicrobial resistance strains from the study area of lactating dairy cows.

2.6.2 Clinical Inspection of the Udder

Udders of the cows were examined by visual inspection and palpation for the presence of any abnormalities. In addition, milk from each quarter was withdrawn and checked for any change in color and consistency (Quinn *et al.*, 2002).

2.6.3 California Mastitis Test (CMT)

The California mastitis test were conducted to diagnose the presence of sub clinical mastitis and it will be carried out according to standard procedures. Squirts of milk from each quarter of the udder were placed in each of four shallow cups in the CMT paddle and an equal amount of the reagent was added. A gentle circular motion was applied in a horizontal plane. Positive samples showed gel formation within a few seconds. The result was scored based on the gel formation and categorized as negative if there was no gel formation, or positive if there was gel formation ranging from +1 to +3 (Appendix 2). If at least one quarter was positive by the CMT then the cow was considered as positive (Quinn *et al.*, 1994).

2.6.4 Culturing procedures

Isolation and identification of *S.aureus* was conducted at Asossa Regional Veterinary Laboratory, on arrival in the laboratory, aliquots (centrifuged milk sample) of 0.01 ml of milk was streaked on blood agar (Oxoid, UK) containing 5-7% sheep blood for isolation of *Staphylococci*. The incubation was done aerobically at 37 °C for 24 hrs. The presence of more than 3 colonies of a similar morph-type was accepted as positive bacteriological finding (Ebrahimi *et al.*, 2010). Identification of the bacteria on primary culture was made on the basis of colony morphology, haemolytic characteristics, Gram stain reaction including shape and arrangements of the bacteria, Catalase test and Oxidase test. In addition, growth characteristics on Mannitol salt agar and purple agar base (1% maltose fermentation) and tube coagulase test was conducted for specifically identifies *Staphylococcus* species (Genta and Heluane, 2001).

2.6.5 Biochemical tests

Isolation and identification of *S. aureus* is done according to standard techniques (Quinn *et al.*, 2002; ISO 6888-2, 2003).

The final identification of the organism and species assignment can be done based on Gram staining, catalase test, carbohydrate dissimilation (mannitol and maltose) fermentation and coagulase test by using rabbit plasma (ISO 6888-2, 2003).

2.6.6 Antibiotic susceptibility test

In determining the type of antibiotic for invitro sensitivity test, retrospective data was compiled on the type of antibiotics used to treat mastitis and other infectious diseases in the region of the study area. In addition to, the selection of the types of antimicrobial agents were made based on clinical considerations including frequent use of the drug in the study area and availability.

The *S. aureus* isolates were tested for anti-microbial susceptibility by disc diffusion method (Quinn *et al.*, 2002). The following antibiotics were used for testing: Cefoxitin (30µg), Vancomycin (30µg), PenicillinG (10u), Tetracycline (30µg), Streptomycin (10µg), Chloramphenicol (30µg), Sulphamethoxazole - trimethoprim (30µg) and Amoxicillin (30µg) Oxoid Company (Hampshire, England).

Colonies isolated from pure culture were transferred into a test tube of 5 ml peptone and suspension was made and incubated at 37°C for 8 hours. The turbidity of the suspension was adjusted comparing with that of 0.5 McFarland

standards. Muller-Hinton Agar plate was prepared and a sterile cotton swab was dipped into the suspension and swabbed on the surfaces of Muller-Hinton Agar plate. Then, the antibiotic discs was placed on the agar plate using sterile forceps and pressed gently to ensure the complete contact with the agar surface. The plates were read after 24 hours of incubation at 35 °C under aerobic condition. The isolates was classified in accordance with the guideline of the National Committee for Clinical Laboratory Standards (CLSI, 2006) as susceptible, intermediate or resistance for each antibiotic tested according to the manufacturer's instructions by measuring the zone of inhibition around the antibiotic disc. Intermediate results were considered as resistant (Huber *et al.*, 2011). Multiple antibiotic resistant (MAR) phenotypes were recorded for isolates showing resistance to three and more antibiotics (Rota *et al.*, 1996).

2.7. Data Management and Analysis

Processing of data was done by computer software. Data was coded and entered to MS Excel spreadsheet and checked for accuracy. After validation, it was transferred and processed using computer software Stata version 12 for analysis. Pearson's chi-square tests was used when appropriate to analyze the proportions of categorical data. Odd ratio and 95% CI was computed, the 95% confidence level was used, and results were considered as significant ($P < 0.05$).

3. RESULTS

3.1. Prevalence of mastitis

In this cross-sectional study, out of the total lactating cows examined, 150(38.96 %) mastitis prevalence was found to be affected with infection. During laboratory examination, 85/385(22.07%) of the *S.aureus* was isolated and 69/385(17.92%) of other coagulase negative staphylococcus spp (CNS) were isolated. The relative proportional prevalence of *Staphylococcus aureus* was 85/150(56.66%). They were found to be statistically significant ($P < 0.00$). The highest mastitic dairy cows distribution were observed in Abrahmo(51.9%) while the lowest prevalence was seen in Ura (29.33%) as indicated in Table 1.

Table 1: Prevalence of mastitic Dairy cows in study sites

Study Sites	N _o of animals examined	Positive	Prevalence (%)	Chi2	p-value
Asossa town	124	42	33.87	11.62	0.009
Abrahamo	104	54	51.9		
Bambasi	82	32	39.02		
Ura	75	22	29.33		
Total	385	150	38.96		

3.2. Risk Factors Associated with mastitis Prevalence

Prevalence of mastitis related to the specific risk factors were determined as the proportion of affected cows out of the total examined. As indicated in (Table 2), the questionnaire survey and observation data result shows previous mastitis history and treatment history, milking hygiene, floor type, and lactation stage, pregnancy status, and blind teat are amongst the potential risk factors, which are associated with mastitis disease in dairy cows farmstead. Accordingly, mastitis prevalence showed significant variation among different blind teat groups ($p = 0.000$), lactation length ($p=0.02$), and pregnancy status ($p=0.004$), previous mastitis history ($p=0.000$) and treatment history ($p=0.000$), milking hygiene ($p=0.005$), floor type ($p=0.01$). However, breed, age, tick infestation, teat lesion and parity have no significant difference with mastitis ($p>0.05$).

Table 2: Result of multivariate logistic regression of attribute risk factors with mastitis

Factor	Categories	Total no examined	No (%) positives	Chi2	p-value
Age(years)	≥3- 5 (y-ad)	137	53 (38.68%)	1.47	0.47
	>6 - ≥9 (adult)	231	88 (38.09%)		
	> 9 (old)	17	9 (52.94%)		
Breed	Cross	180	70(38.88%)	0.00	0.97
	Zebu	205	80(39.02%)		
Parity	1-2	209	80(38.27%)	2.83	0.24
	3-4	123	44(35.77%)		
	≥5	53	26(49.05%)		
Lactation Stage (m)	Early (≤3)	124	60(48.38%)	10.42	0.02
	Mid (4-6)	139	41(29.49%)		
	Late (7-9)	84	32(38.09%)		
	Dry (>9)	38	17(44.7%)		
Pregnancy Status	Pregnant	103	28(27.2%)	8.20	0.004
	Non- Pregnant	282	122(43.26%)		
Previous mastitis History	Infected	141	135(95.74%)	301.6	0.000
	Non- infected	244	15(6.14%)		
Floor type	Concrete	263	91(34.6%)	6.63	0.01
	Muddy (soil)	122	59(48.36%)		
Milking hygiene	Good	270	93(34.44%)	7.75	0.005
	Poor	115	57(49.56%)		
Prevoius mastitis Rx history	Yes	96	92(95.83%)	173.9	0.000
	No	289	58(20.06%)		
Blind teat	No	331	96(29.0%)	40.09	0.000
	Yes	54	54(100%)		
Tick infestation	No	331	134(40.48%)	2.29	0.13
	Yes	54	16(29.62%)		
Teat lesion	No	366	146(39.89%)	2.69	0.10
	Yes	19	4(21.05%)		

3.3. Antimicrobial Susceptibility Test

Antimicrobial susceptibility tests were performed on 22 staphylococcus isolates and were tested for antimicrobial sensitivity for 5 different types of antibiotics. The present study has demonstrated the existence of the levels of resistance of *S.aureus* to commonly used antimicrobial agents. 76.92 % of the *S. aureus* was found to be resistance to Cefoxitin. The resistance profile of Amoxicillin, Penicillin G, Tetracycline, and Gentamycin, were 84.61, 78.84%, 69.23%, and 53.84%, respectively (Table-3). In this study, *S. aureus* were found to be highly susceptible to Cloxacillin (63.46%) followed by Gentamycin (40.38%). However, these isolates were highly resistant to penicillin G (78.84%) and Cefoxitin (76.92%) followed by Tetracycline (69.23%). The antimicrobial resistance profiles are shown in Table 3.

Table 3: Resistance and susceptible of *S. aureus* isolates to different antimicrobials (n = 22).

Antimicrobial agents	Disc content (µg)	No. of Isolates	Resistance	Intermediate	Susceptible
			No_(%)	No_(%)	No_(%)
Cefoxitin	30	22	17(77.27)	0	5(22.72)
TTC	30	22	15(68.2)	2(9.09)	5(22.72)
Cloxacillin	5	22	5(22.72)	3(13.63)	14(63.63)
Gentamycin	10	22	12(54.54)	1(4.54)	9(40.90)
Penicillin G	10	22	17(77.27)	0	5(22.72)
Mean			66 (13.2)	6 (1.2)	39(7.8)

Key: S- Susceptible, I- Intermediate, R- Resistant

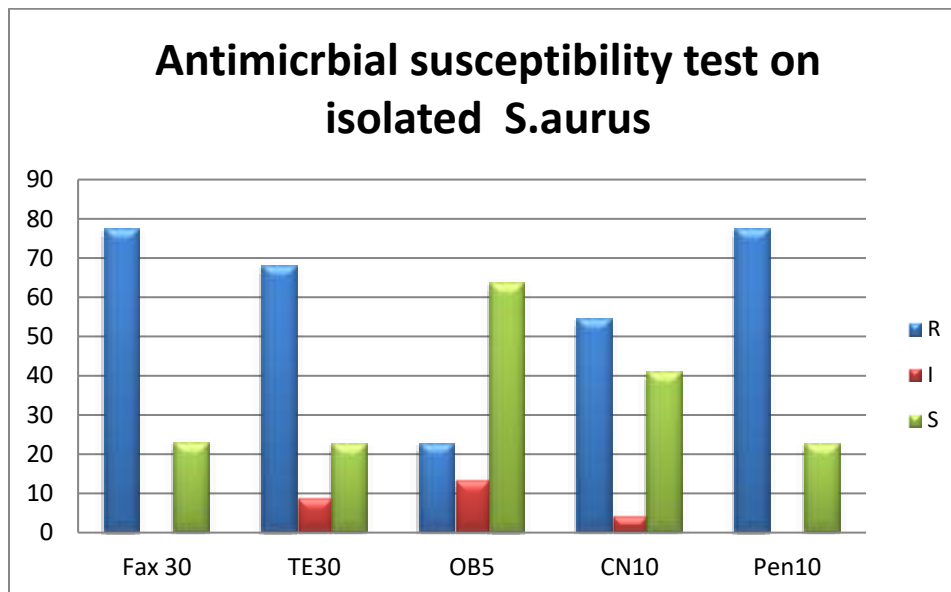


Figure 4: Antimicrobial drugs resistance and susceptibility profile of *S.aureus* isolated from milk
Key: TE30=Tetracycline, Fax30= Cefoxitin, CN10=Gentamycin, Pen10= Penicillin G, OB5= Cloxacillin.

4. DISCUSSION

In the present study, the overall prevalence of mastitic dairy cows were 38.96 % in cows. This result was in line with the earlier reports by Biniam (2014) in and around Wolaita Sodo, Abinet (2015) in and around Batu town, Kerro and Tareke (2003) in Southern Ethiopia, (40.9%), (42.59%), (40%) in cows respectively. This report is relatively similar with the assertion by Radostits *et al.* (2000) that, in most countries and irrespective of the cause, the prevalence of mastitis is about 50% in cows and 25% in quarters. Besides, this result was in line with the findings of Bitew *et al.* (2010) at Bahir Dar, and Mulugeta and Wassie (2013), around Wolaita Sodo, 28.8%, 29.5% in cows respectively.

However, this finding is lower when compared with the previous findings of Shimelis (2014) in Selale/Fitche area, Alemayehu (2015) in Bahir Dar and its surroundings, Mesfin (2015) in and

around Kombolcha, (83.1% v 65.42%), (62.06% v 42.44%), (56% v 33.7%) in cows and quarters respectively. In addition, it disagrees with the previous findings of Sori *et al.* (2005) in and around Sebeta, Lakew *et al.* (2009) in Asella, Abaineh (1997) in Fiche, Abera *et al.* (2013) in Adama, Zerihun (1996) in Addis Ababa, Mekibib *et al.* (2010) in Holeta, Nesru (1986) in Dire-Dawa, 52.78%, 64.4%, 65%, 66.6%, 68.1%, 71.0%, 85.6% in cows respectively. This variability in prevalence of mastitis between different reports could be attributed to differences in farms management practice or to differences in study methods agro-climatic condition. As mastitis is a complex disease involving interactions of various factors such as managerial and husbandry, environmental conditions, animal risk factors, and causative agents, its prevalence will vary (Radostitis *et al.*, 2007).

With regard to the bacteriological analysis of milk sample, the relative isolates of *S.aureus* were 85/150(56.66%). This finding is inconsistent with the earlier findings of (51.56%) by Shimelis (2014), in Selale /Fiche Area, around Sebeta (44.03%) by Sori *et al.* (2005), in Holleta agricultural research centre (43.3%) by Duguma *et al.* (2013), in Hawassa area (48.75%) by Daka *et al.* (2012), in Holeta town (47.1%) by Mekibib *et al.* (2010) and in Debre Ziet area (39.5%) by Addis *et al.* (2011). Similarly, this result was inline with the previous findings of Bedada and Hiko, (2011), Workineh *et al.* (2002) and kerro and Tareke, (2003) who have reported as 39.1%, 39.2% and 40.3% *S. aureus* isolates at Assela, Addis Ababa and Southern Ethiopia, respectively. It was also closely comparable with findings of Lakew *et al.* (2009) and Ndegwa *et al.* (2000) who reported 41.1% and 43.3% in dairy cows, respectively.

However, *S. aureus* isolate is high as compared to the previous findings of Mesfin (2015) in Kombolcha, Abinet (2015), in Batu, Abebe *et al.* (2013), in Addis Ababa, by Seedy *et al.* (2010) in Egypt, Biniam (2014) in Wolta Sodo, Alemayehu (2015) in Bahir Dar, Hussein *et al.* (1997), Bishi (1998) and Mekuria *et al.* (2013), 26.7%, 17.13%, 16.0%, 17.2%, 18.39%, 15.02%, 10 %, 9%, 16% respectively. The high prevalence of this organism may be associated with its frequent colonization of teats, its ability to exist intracellular and localize within micro abscesses in the udder and hence resistant to antibiotic treatment (MacDonald, 1997). The Bacteria usually establish chronic, subclinical infections and are shed in the milk, which serves as a source of infection for other healthy cows during the milking process. The possible explanation for the variation might be that *S. aureus* is a contagious pathogen transmitted from one cow to another or individual by contact with animals during unhygienic milking procedures (Rowe, 1999). Therefore, the *S.aureus* occurrence at a considerable high percentage indicates the alarming situation for dairy farms.

The prevalence of mastitis in local zebu and cross breeds were in-significantly associated with the occurrence of mastitis ($p>0.05$). Comparable research works were reported by Almaw *et al.* (2009) in Gondar town and its surroundings, Sori *et al.* (2005) in and around Sebeta showed that breed significantly influenced the occurrence of mastitis.

In addition, this finding was closely similar with Bitew *et al.* (2010) who reported in Bahir Dar, between Cross and Fogera breed, Lakew *et al.* (2009) in cross and local Arsi breed and Biffa *et al.* (2005) found significant difference between local Zebu, Holstein-Frisian and Jersey breeds in Ethiopia, That was Holstein Fresian pure breeds were affected at a

higher rate both by clinical (26.3%) and subclinical (30.1%) mastitis than local breeds. Increased milk yield from genetic selection may be accompanied in genetic susceptibility to mastitis (Schutz, 1994). Besides this, the low occurrence of mastitis in local breeds in addition to genetic factors could also be one indication for higher occurrence of mastitis prevalence in areas where exotic breeds and their hybrids well adapted. Therefore, the lower prevalence in local zebu breeds in this study could be associated with difference in genetically controlled physical barrier like streak canal sphincter muscles, keratin in the teat canal or shape of teat end where pointed teat ends are prone to lesion (Seykora and Mcdaniel, 1985). In addition to physical barriers, the difference in occurrence of mastitis in these breeds could arise from differences in cellular immunity (Erskine, 2001).

The observed higher occurrence of mastitis during early lactation as compared to mid and late lactation stages was significant ($p<0.02$). The finding of higher infection in cows in early lactation stage followed by late and medium lactation stages in the study concurs with previous reports of Mulugeta and Wassie, (2013); Biffa *et al.* (2005) and Tamirat, (2007). In cows most new infections occur during the early part of the dry period and in the first two months of lactation (Radostits *et al.*, 2007). This may be due to an absence of dry period therapy and birth related influences. During a dry period, due to low bactericidal and bacteriostatic qualities of milk, the pathogens can easily penetrate into the teat canal and multiply (Aylate *et al.*, 2013). Radostits *et al.* (2000) suggested that, the mammary gland is more susceptible to new infection during the early and late dry period, which may be due to the absence of udder washing and teat dipping, which in turn may have increased the presence of potential pathogens on the skin of the teat. Moreover, during a dry period due to the low bactericidal and bacteriostatic qualities of milk, the pathogens can easily penetrate into the teat canal and multiply; this can be carried over into the post parturient period and ultimately develop into mastitis.

In this study, floor system had a significant influence on the occurrence of mastitis ($p=01$). In agreement with Abera *et al.* (2013) in Adama town and Fekadu *et al.* (2005) in southern Ethiopia, Lakew *et al.* (2009) and Sori *et al.* (2005). The findings of a high prevalence of mastitis in farms with muddy (soil) floors (48.36%) when compared with concrete floor types (35.22%) shows the occurrence of mastitis is significantly associated with the housing (bedding) type or condition of the farm. This is due to association with poor sanitation and cows which were maintained in dirty and muddy common barns with bedding materials that favor the proliferation and transmission

of mastitis pathogens. The main sources of infection are udder of infected cows transferred via milker's hand, towels and environment (Radostitis *et al.*, 2007).

Occurrence of mastitis was significantly associated with milking hygienic practice ($p=0.005$). Cows at farms with poor milking hygiene standard are severely affected (49.56%) than those with good milking hygiene practices (35.03%) (Mulugeta and Wassie, 2013; Lakew *et al.*, 2009; Sori *et al.*, 2005). This might be due to absence of udder washing, milking of cows with common milkers' and using of common udder cloths, which could be vectors of spread especially for contagious mastitis (Radostitis *et al.*, 2007).

In this finding the prevalence of mastitis was not significantly influenced by age categories ($P > 0.05$). Similar result was reported by Shimelis (2014) in Selale /Fiche, no significant effect ($p > 0.05$). In this study, parity is not significantly influenced on the occurrence of mastitis ($p > 0.05$). In contrast to this study, the increased occurrence of mastitis with parity was reported by Mekibib *et al.* (2010) in Holeta town and Haftu *et al.* (2012) in northern Ethiopia.

The present study showed that the resistance of *S. aureus* to Penicillin G (78.84%), Cefoxitin (76.92%), Tetracycline (69.23%), Cloxacillin (23.07%) and Gentamycin (53.84%) observed in milk samples. Comparable research works were reported in various parts of Ethiopia by Biniam T (2014) revealed resistance of *S. aureus* to Penicillin G (100%), Cefoxitin (71.8%), and Tetracycline (69.2%) in and around Wolaita Sodo, southern, Ethiopia. Besides this, Alemayehu (2015) indicated resistance of *S. aureus* to Penicillin G (95.8%), Cefoxitin (75.7%), and Tetracycline (72.2%), from Bovine mastitic milk in Dairy farms of Bahir Dar.

In addition, this research is in accordance with the findings of Abebe *et al.* (2013) who reported resistant of *S. aureus* to penicillin G 96.7% and tetracycline 73.8% around Addis Ababa, and Abera *et al.* (2010) 94.4% resistance to penicillin G in Adama; in addition to this study has demonstrated the existence of alarming level of resistance of *S. aureus* to commonly used antimicrobials (penicillin G, and tetracycline) in dairy farms. This results were in consistent with reports from earlier studies in the other countries (Edward *et al.*, 2002; Gentilini *et al.*, 2002 and Jakee *et al.*, 2008) suggesting a possible development of resistance from prolonged and indiscriminate usage of some antimicrobials. Hence, penicillin and tetracycline are the only most commonly used antimicrobials for the treatment of other infections as well as mastitis in veterinary practice in Ethiopia, as the result, there was spread of

drug resistance reported by many researchers which was in line with the recent findings.

The resistance of *S. aureus* isolates to beta-lactam antibiotic was evident. High percentage of *S. aureus* was resistant to the most frequent drugs. In agreement with the finding of by Derese *et al.* (2012), the study showed cefoxitin resistant isolates were obtained from the milk. All cefoxitin resistant *S. aureus* were also resistant to penicillin G. Out of the (76.92%) cefoxitin resistant *S. aureus* isolates, (84.61%) and (78.84%) were also resistant to amoxicillin and Penicillin G respectively. This is an indicator of MRSA (Daka *et al.*, 2012). This is due to the fact that resistance of *S. aureus* to these drugs may be attributed to the production of β -lactamase, an enzyme that inactivates penicillin and closely related antimicrobials (Wubishet *et al.*, 2012; Sharma *et al.*, 2011; Green and Bradely, 2004).

In the present observation, frequent multidrug resistance pattern were exhibited for Penicillin G, Cefoxitin and tetracycline. Comparably, Alemayehu (2015) who reported as resistance for multidrugs, mainly to penicillin G, Cefoxitin and tetracycline. In addition, Shimelis (2014) who found that, 86.46 % of the isolates were resistant to different combinations of two or above tested antibiotics and the most frequent multidrug resistance pattern consisting of three drugs' is exhibited for, gentamicin, ceftazidime and streptomycin with a resistance of 9.46% of the isolates. Similar finding by Mekuria *et al.* (2013) reported MRSA isolate with resistant to more than two of non- β -lactam antimicrobials. This multi drug resistance occurred might be due to administration of multiple antibiotics for prophylaxis or infection, lack of drug sensitivity tests in the dairy farms, uncontrolled or discriminate use of antibiotics in the farms and another possibility is that cattle are being treated with antibiotics for other conditions, thereby selecting for resistant populations of *S. aureus* (Shitandi and Sternesjo, 2004).

5. CONCLUSION AND RECOMMENDATIONS

Dairy cows mastitis could be one of the major constraints to dairy production in extensive dairy farms. Different potential risk factors were associated with mastitis in the study area, amongst these, length of lactation, blind teat, milking hygiene, floor type, lactation stage, previous mastitis & treatment history and pregnancy status of the animal were prominent. Mastitis caused by *S. aureus* at cow was one of the major problems of dairy cows in milk production. It was found that the majority of the tested isolates were resistant to the various antimicrobial agents especially penicillin G, Cefoxitin, and Tetracycline.

Based on the above conclusion the following points are forwarded:-

- Mastitis control strategy should be initiated and promoted in the study area;
- Hygiene measures during milking procedure should be practiced that may reduce the transmission of the disease
- There should be regular antimicrobial sensitivity test to select effective and alteration of antibiotics to reduce the problems of drug resistance development towards commonly used antibiotics

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