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SOME BACTERIOLOGICAL STUDIES ON SUB CLINICAL MASTITIS IN CATTLE AND ITS RELATION TO CHANGES IN THE MILK PROTEIN ELECTROPHORETIC PATTERN

(With 7 Tables)

By

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**بعض الدراسات البكتريولوجية لمرض التهاب الضرع الخفى فى الأبقار وعلاقته
بالفصل الكهربى لبروتينات اللبن**

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أجريت هذه الدراسة على عدد 50 بقرة حلابة تم إختيارها عشوائيا من مختلف المزارع الصغيرة بمدينة أسيوط حيث تم تجميع عدد 180 عينة لبن من هذه الأبقار وذلك لفحصها معمليا اولاً بإختبار الكاليفورنيا لتحديد المصاب منها بالتهاب الضرع الخفى ثم فحص الإيجابى لهذا الإختبار بكتريولوجيا لعزل وتصنيف الميكروبات المسببة. وقد وجد ان 58 عينة من هذه العينات كانت ايجابية لإختبار الكاليفورنيا بنسبة 32.2%، ثم بفحص هذه العينات بكتريولوجيا وجد أنها إيجابية بنفس النسبة 32.2%. هذا وقد تم عزل عدد 102 عترة تمثل مجموعتين من المسببات البكتيرية الأولى عالية الضراوة وتمثل عدد 75 عترة بنسبة 73.5% أما المجموعة الثانية فتمثل ميكروبات التلوث البيئى 27 عترة بنسبة 26.5%، وقد تم عزل أكثر من عترة من الميكروبات المرضية من العينة الواحدة من عدد 28 عينة لبن بنسبة 48.3%. وتمثلت العترات المعزولة عالية الضراوة فى عزل الميكروب العنقودى الذهبى بنسبة 29.4% والميكروب العنقودى السالب للتجلط بنسبة 39.3%، والميكروب السبحى اجلاكتيا بنسبة 4.9%. أما ميكروبات التلوث البيئى فقد تمثلت فى الميكروب القولونى 9.8%، والانتيروكوكاس والكليبيسيلا والباسيلس سيريس بنسبة 7.8%، 1.9%، 6.9% على التوالى. وقد تم عمل اختبار حساسية لهذه العترات المعزولة لاختيار المضادات الحيوية الأنسب فى العلاج، كذلك ناقش البحث الأهمية الوبائية والصحية والاقتصادية لمرض التهاب الضرع الخفى فى الأبقار والإجراءات الواجب اتباعها لمنع إنتشار المرض بين حيوانات المزرعة. كما لوحظ من خلال الفصل الكهربى لبروتينات اللبن فى حالات إتهاب الضرع الخفى فى الأبقار إرتفاع معنوى إحصائى فى مستوى البروتين الكلى فى مصل اللبن وفى الأجسام المناعية والألبومين باللبن بينما شهد كل من الألفا والبيتا لاكتو ألبومين نقص معنوى حاد كما صاحب حالات الإتهاب الضرع الخفى أيضا انخفاض معنوى إحصائى فى كلا من الألفا كازين والبيتا كازين والكابازين.

Key words: *Sub clinical mastitis, bacteriology, milk protein, electrophoresis.*

SUMMARY

This study was conducted on 50 dairy cows, which were randomly collected from different smallholder farms in Assiut city. 180 milk samples representing 50 dairy cows under test were firstly screened for the presence of subclinical mastitis by using California Mastitis Test (CMT); 58 milk samples yield positive results, and all of these were positive for the bacteriological examination. The isolated bacterial strains (102 isolates) including contagious bacteria representing 75 strains (73.5%) and environmental bacteria 27 (26.5%). 48.3% of milk samples showing mixed infection. The isolated contagious strains were *Staph aureus* (30 strain), coagulase negative *Staph* (CNS) 40 strain, *Strept agalactia* 5, while the environmental bacteria were *E coli* 10, *Enterococcus sp* 8, *Klebsiella spp* 2 and *Bacillus spp* 7. The electrophoretic pattern of milk proteins showed that there were significant statistical increase in total whey protein, immunoglobulin and albumin in cows' milk samples with sub clinical mastitis. While α -lactalbumin, and β -lactoglobulin showed significant decreased. There were a highly significant decrease in α -casein and κ casein in sub clinical mastitic samples in comparing with the normal milk. In addition, there were a significant decrease in β -casein and non-significant decrease in γ casein milk protein fractions in sub clinical mastitic milk. Recommendations were suggested to eradicate and control subclinical mastitis to decrease the economic losses in lactating dairy cows.

INTRODUCTION

Bovine mastitis is a complex and economically important infectious disease for dairy cattle throughout the world, which can result in substantial losses due to reduced milk yield, and increase culling rates and veterinary expenses (Miles *et al.*, 1992).

Subclinical mastitis, without any signs of inflammation compared with clinical mastitis, is the main form of the disease, and accounts for the majority of bovine mastitis cases in dairy herds (Oliver *et al.*, 2004)

Sub-clinical mastitis is 3-40 times more common than the clinical mastitis and causes the greatest overall losses in most dairy herds (Schultz *et al.*, 1978).

The identification of subclinically mastitic quarters and cows is of great concern in the control programs to face the spread of infection within and between dairy herds. Li, *et al.* (2009) found that 28% of tested quarters having subclinical mastitis.

The prevalence of subclinical mastitis is known to be influenced by many additional factors, such as husbandry, management, genetics, nutrition and associated metabolic and endocrine changes (Wielgosz-Groth and Groth, 2003).

More attention has been given for the diagnosis of SCM by indirect test (Joshi, *et al.*, 1976). Bacteriological culture is a test, similar to CMT and somatic cell count (SCC), which are extremely useful tools for monitoring udder health status and for problem- solving (Radostits, *et al.*, 1994). Simply the bacteriological culture of milk is essential to determine the presence and type of pathogens involved in cases of intra-mammary infection (IMI). Bacterial pathogens that cause mastitis are generally classified as either contagious or environmental based upon their primary reservoir and mode of transmission (Makovec and Ruegg 2003). Contagious bacteria, such as *Staphylococcus aureus*, *Streptococcus agalactia*, and *Corynebacterium bovis*, can spread from an infected cow to another one (Abdel- Khalek and Sherbini, 2005). While environmental bacteria, such as: *E.coli*, *Strept uberis*, *Actinomyces pyogenes*, *Pseudomonas aeruginosa* and other *Staphylococcus spp*, are commonly present in surrounding environment and may reach the teat end from that source, (Anwer, *et al.*, 2003).

Staphylococcus aureus seemed to be the predominant organism causing subclinical mastitis (Kader *et al.*, 2002). It may predispose the herd for infection by coliform or other pathogens (Ibtisam *et al.*, 1993). *Streptococcus agalactiae* lives in milk and in the mammary gland, but can survive only for a few hours outside of the mammary gland (Phuektes, *et al.*, 2001).

Subclinical mastitis caused by intramammary infection (IMI) with coagulase-negative staphylococci (CNS) is common in dairy cows and may cause herd problems (Thorberg *et al.*, 2009).

The identification of subclinically mastitic causing pathogens and their antimicrobial sensitivity testing are important points in the implementation of control programs (Dhakal *et al.*, 2007).

Detection and evaluation of proteins in milk during the course of mastitis are important to elucidate the pathologic mechanism of bovine mastitis (Kato *et al.*, 1989). Electrophoresis has played an important role in the study of milk proteins and has been an integral part of research on the genetic variants of the major proteins components of milk. Indeed, the designations of caseins are derived from electrophoretic analysis and minor caseins components were discovered by electrophoresis (Kostyra, 1990).

Bovine milk contains 3.0-3.5 % (w/v) protein (Cole, 1986). Caseins are 2.4-2.8% of fluid milk and exist in milk as a micelle containing the four casein types: α 1- α 2- β and k-casein (Kostyra, 1990). Whey proteins are serum albumin, α -lactalbumin, β Lactoglobulin, and immunoglobulin.

Microbial toxins and enzymes from damaged cells cause injury of secretory cells (Kitchen, 1981). Therefore, the ability of the mammary epithelium to synthesize and secrete the major specific milk constituents is reduced (Fox *et al.*, 1985 and Eberhart *et al.*, 1987). While the secretion of other proteins like lactoferrin is simultaneously upregulated (Schmitz *et al.*, 2004). The concentration of caseins is reduced in infected quarters due to reduced secretion and due to destruction by blood-borne proteases like plasmin (Politis *et al.*, 1992).

Aim of the work: To study the prevalence of subclinical mastitis between cattle in Assiut governorate and to determine the most prevalent causative agents, also studying the relationship between different bacterial causes of subclinical mastitis and the changes occur in milk protein fractions.

MATERIALS and METHODS

Animals:

A total of 50 dairy cows, were randomly selected from different smallholder farms at Assiut governorate. The udder and teats of the selected cows were physically and clinically investigated to exclude the clinical mastitis, all animals in all farms were hand milked, twice daily.

Milk Samples:

Udders and teats of the selected cows were thoroughly washed and dried with sterile clean towel and disinfected with 70 % alcohol. The first milking streams were rejected and 20 ml of milk / each quarter was collected in sterile screw capped bottle. The collected samples were divided into three portions, one portion was subjected to California Mastitis test (CMT), another was subjected to bacteriological examination and about 10 ml of milk / each quarter were used for milk separation into milk serum (whey) and Casein.

Screening test:

A total of 180 milk samples were subjected to California Mastitis test (CMT) according to Schalm *et al.* (1971) for detection of sub clinical mastitis. CMT scored from 1 to 5 corresponding to no reaction, trace, mild reaction, moderate reaction and strong reaction, respectively. The positive samples were subjected to bacteriological examination and electrophoresis of milk protein.

Bacteriological Examination:

The milk samples were incubated for 18-24 hours at 37 °C, and 10 ml of milk samples were transferred into sterile small centrifuge tubes. The tubes were centrifuged at 3000 rpm for 20 minutes, and then the cream and the supernatant were discarded to obtain sediment, then a loop full from milk sediment was streaked onto Azid blood agar plate (Cruickshank, *et al.*, 1975) and a loopfull also was inoculated into nutrient broth (Diffco), Na Cl broth 10%, MacConkey broth (Oxide). The previously inoculated tubes were incubated at 37 °C for 24 hours. From the incubated tubes, loopfulls were streaked onto the surface of the nutrient agar, blood agar with 5% sheep blood, MacConky agar (Oxoid). The inoculated plates were incubated aerobically at 37°C for 24 h.

The suspected colonies were identified morphologically by Gram's stain and biochemically confirmed according to Quinn, *et al.* (1994), using catalase and coagulase tests. Identification of *Streptococci spp* was done by catalase test, hemolytic activity, sodium hippurate hydrolysis, aesculin hydrolysis on blood agar with 0.1% aesculin. *Enterobacteriace* were identified biochemically by conventional IMVIC (Indole, Methyl red, Voges Proskauer and citrate utilization) test, Motility, triple sugar iron agar (TST) inoculation, according to (Quinn, *et al.*, 1994; DeBoer and Heuvelink, 2000).

Antibiotic sensitivity testing:

Antimicrobial susceptibility testing by using disc diffusion standard technique according to Bauer *et al.* (1966) and Finegold and Martin (1982) was applied. The isolated micro-organisms were tested against Penicillin 10 i.u., Amoxicillin/Clavulanic acid 20/10 mcgm, Chloramephenicol 30 mcgm, Gentamicin 10 mcgm, Norfloxacin 10 mcgm.

Sample preparation for electrophoresis:

Fresh warm raw milk was obtained from Holstein cows. Within an hour after milking, milk samples were skimmed by centrifugation at 3000 rpm for 15 min to remove their creams and cells. Samples were then treated with 0.1 M, hydrochloric acid at the controlled PH of 4.8 for casein precipitation. Treated samples were recentrifuged and the supernatants (Whey) were collected. The casein precipitate was separated from the whey by filtration. The casein fraction was washed with distilled water three times, maintaining the PH of the water at 4.7 by addition of dilute HCl.

Casein solution was prepared by dissolving 1g casein in 100 ml of 100 mM Tris-Hcl buffer (PH 8.0) and heating in a boiling bath for 20 min. The solution was filtrated without cooling and stored at 4 °C until to be used. Casein solutions must be discarded after 2 days. The total whey

protein was determined according to Henry, (1969), application of SDS-polyacrylamide gel electrophoresis according to Sambrook, *et al.* (1989). The protein standard is ranged from (10-250KDa) BioRad, using methods of Sambrook, *et al.* (1989).

Statistical Analysis:

Student's t-test was carried out to find the differences between the results of mastitic and non mastitic milk samples and the results were given as mean \pm SEM.

RESULTS

Table 1: Udder-quarter level prevalence of subclinical mastitis:

No. of Quarters	CMT positive		Bacteriologically positive		CMT negative		Bacteriologically negative	
	No.	%	No.	%	No.	%	No.	%
180	58	32.2	58	32.2	122	67.8	122	67.8

Table 2: Prevalence of sub clinical mastitis by using CMT:

No. of cows	CMT positive		CMT negative	
	No.	%	No.	%
50	27	54	23	46

Table 3: Frequency percentages of single and mixed infection in quarter milk cow's samples

	Single infection	Double infections	Triple infections
No.	30	18	10
%	51.7	31	17.3

Table 4: Prevalence of the isolated subclinical mastitis bacteria from the examined cow's milk samples:

Isolated species	No.	%
<i>Staph aureus</i>	30	29.4
<i>Coagulase Negative Staph (CNS)</i>	40	39.3
<i>Streptagalactia</i>	5	4.9
<i>E coli</i>	10	9.8
<i>Enterococcus spp</i>	8	7.8
<i>Klebsiella spp</i>	2	1.9
<i>Bacillus spp</i>	7	6.9
Total	102	100

Table 5: Results of Antibiotic sensitivity testing of the isolated strains

	Penicillin	Amoxicillin/ Clavulanic acid	Chloramphenicol	Gentamicin	Norfloxacin
Staph aureus	50%	50%	100%	75%	100%
CNS	0%	25%	100%	100%	100%
Streptagalactia	100%	100%	100%	100%	100%
E coli	0%	0%	100%	50%	100%
Klebsiella spp	0%	0%	100%	100%	100%
Bacillus spp	0%	0%	100%	100%	100%
enterobacter	0%	0%	100%	0%	100%

N.B.: the antimicrobial agents tested were selected on the basis of the actual veterinary practice.

In this study, there were no differences between the different bacterial infections (which causing subclinical infections) in the whey or casein milk protein electrophoretic pattern or even the mixed infection.

The percent of protein fractions were significantly different between normal and all bacterial infection ($p < 0.01$), but there were no differences between different kinds of bacteria so in this study, we comparing between normal and mixed infection causing sub clinical mastitic milk.

In case of sub clinical mastitis, total whey protein significantly increased ($P < 0.05$) and was accompanied by significant increase ($p < 0.01$) in the immunoglobulin and albumin content in milk. While both α – lactalbumin, and β -lactoglobulin were significantly decreased (Table 7).

In this study, there were a highly significant decrease ($P < 0.01$) in α -casein and κ casein in sub clinical mastitis in comparing with the normal milk. Also there were a significant decrease in β -casein and non significant decrease in γ casein milk protein fraction in sub clinical mastitis milk (Table 8).

Table 6: Percentage (%) of whey-protein fractions in normal and subclinical mastitic milk:

Protein fraction	Normal milk	Sub clinical Mastatic milk
Total whey protein	14.77±1.34	17.68±2.20*
Immunoglobulin	14.91±1.10	19.6±1.43**
Albumin	7.95±0.54	15.10±1.27**
α –lactalbumin	26.72±0.45	22.25±0.39**
β -lactoglobulin	54.03±0.64	31.00±1.02**

*significant at $P < 0.05$., ** Significant at $P < 0.01$.

Table 7: The percentage (%) of casein-protein fractions in normal and subclinical mastitic milk.

Protein fraction	Normal milk	Sub clinical Mastatic milk
α - casein	10.68±0.21	6.26±0.25**
β -casein	8.89±0.20	7.27±0.19*
κ casein	16.26±0.09	14.22±0.18**
γ casein	22.20±0.21	21.48±0.20

*significant at $P < 0.05$., **Significant at $P < 0.01$ and

DISCUSSION

Cows suffering from subclinical mastitis show no signs, secret apparently normal milk for long time during which infected animal act as potential reservoir for the responsible causative organisms and spread infection among neighboring animals in the herd (Mohamed *et al.*, 1993).

The California mastitis test (CMT) is used on farms to identify sub clinical mastitis by an indirect estimation of the somatic cell count (SCC) in milk.

Results in Table (1) revealed that the quarter level prevalence of sub clinical mastitis in cows based on the results of CMT and bacteriological examination were 32.2 % (58 out of 180), these results were completely agree with that recorded by Hawari and Al-Dabbas (2008) as they recorded that 31.4% of tested quarters showed subclinical mastitis, and somewhat similar to the results recorded by Sadek (2008) (29.55%) and Li *et al.* (2009) (28%).

Table (2) estimates the animal prevalence of subclinical mastitis in cow's milk samples based on the results of CMT. Out of 50 cows examined 27 animals (54%) gave positive results. Similar results were detected by Li *et al.* (2009) as they recorded that 54.3% of tested cows were sub clinically mastitic cases and by Tijare *et al.* (1999) (57.98%). A lower percentage was estimated by El-Kholy and Hosein (1990) 16.6%, while higher percentages were recorded by Sadek (2008) 59.05% and Sexena *et al.* (1993) 64%.

The quarter prevalence of subclinical mastitis in cow's milk samples based on the results of bacteriological examinations was illustrated in Table (1). Out of 180 quarter cow's milk samples examined, 58 (32.2%) were positive.

These findings were lower than that recorded by Sharma and Rai (1977), Hatem *et al.* (1984), Saini *et al.* (1994), Ismail and Hatem (1998) and Nazem and Azab (1998) as they recorded 40.4, 67.74, 76.13, 87.5 and 75.25% respectively, while these results are higher than that recorded by Ismail and Hatem (1998) 26.5% and Sadek (2008) 28.5%. The subclinical mastitis incidence varied widely due to changing management conditions (Radostitis *et al.*, 2000).

CMT field test is a dependable and a reliable perfect test (El-Gamal, 1989 and El-Balkemy *et al.*, 1997). In this study CMT results were agree completely (100%) with bacteriological isolation.

Table (3) illustrated that the mixed infection was 48.3% and the single isolation 51.7%, this finding reflects an idea about the level of environmental bacterial contamination in the herd and demonstrate the complexity of the disease.

Table (4) shows the isolated bacterial strains from the examined subclinical mastitic milk samples, it is revealed that the main isolates were of the contagious type *Staph aureus*, *CNS*, *Strept agalactia* in percentages of 29.41%, 39.21%, and 4.9% respectively. The contagious organisms were well adapted to survive in the udder and usually establish mild subclinical infection for long duration (El-Khodery and Hoedemaker, 2005 and Abdel-

Khalek and El-Sherbini, 2005) and can spread from infected quarter to another quarters (El-Balkemy *et al.*, 1997). Staphylococci typically colonize the broken skin and can enter the udder through abrasions of the teat (Dhakal, 1997).

Several studies have estimated the prevalence of subclinical mastitis due to *Staph aureus* and have shown wide variation. The obtained results 29.41% were similar to that recorded by Janosi and Balty, 2004, and lower than that recorded by (Attia *et al.*, 2003) (80%) and Shitandi and Kihumbu, 2004 (45.6%). *Staph aureus* and *Streptagalactia* are commonly isolated from sub-clinical mastitis (Abdel-Khalek and El-Sherbini, 2005) where *Staph aureus* commonly produce long-lasting infections as it developed sophisticated system to avoid phagocytosis or macrophages (Vanfurth and Van Zwet, 1986).

The importance of coagulase negative Staph as a cause of sub-clinical mastitis was previously demonstrated by Hodges *et al.* (1984), Malinowski *et al.* (1992).

Subclinical mastitis caused by intra-mammary infections (IMI) with coagulase-negative staphylococci (CNS) is common in dairy cows and may cause herd problems. Control of CNS mastitis is complicated by the fact that CNS contains a large number of different species (Shitandi and Kihumbu, 2004).

From Table (4) it is clear that the CNS were isolated from 39.3% of subclinically mastitic milk samples and this completely agree with Sargeant *et al.* (2001) who isolate CNS from 40% of tested samples.

Table (4) illustrate the environmental bacteria (*E coli*, *enterococcus spp*, *Klebsiella spp*, and *Bacillus spp*) isolated from the subclinically mastitic quarters cows milk samples, these environmental bacteria originate from the surrounding environment including air, soil, water, bedding materials, faecal matter, milking man, and milking utensils (Anwer *et al.*, 2003). The portal of entry into mammary gland for gram negative bacteria is the teat canal.

The most familiar environmental pathogen, *E.coli* is widely documented to be a sub-clinical mastitis pathogen (Anwer *et al.*, 2003). Its persistence within the mammary environment was of the recurrent quarter's *E.coli* mastitis and its spread among other quarters and cows may occur during the milking process (Bradley and Green, 2001).

Table (5) showing the results of antimicrobial sensitivity testing, it is clear that the most effective antimicrobial agent are Norfloxacin and chloramephnicol followed by Gentamicin and the least effective one is the penicillin and this is may be attributed to the misuse of penicillin in the

veterinary practice (incomplete treatment course and under dosing) so resistant strains were developed.

Inflammation of the mammary gland leads to a variety of compositional changes in milk either because of local effects or because of serum components entering the milk and the movement of some normal milk components out of the alveolar lumen into the perivascular space (Harmon, 1994).

Albumin content of milk in sub clinical mastitis was significantly increased compared to the healthy ones. The increase of albumin content during mastitis has been reported in cows (Coulon *et al.*, 2002 and Batavani *et al.*, 2007).

De Wit, 1998, said that the main site of albumin synthesis is in the liver, and the albumin enters the milk by leaking through the epithelial tight junction from the blood stream, while Shamay *et al.* (2005) found that the extra hepatic synthesis of albumin has been demonstrated in mammary gland epithelial cells, but in lesser amounts than the liver.

The marked increases of albumin in mastitic cows suggest that a major source of the increase in the content of albumin in milk under inflammatory conditions is the mammary gland itself.

Immunoglobulin in mammary secretions is serum- derived or produced in the udder and pass into the milk through the mammary epithelium. The concentrations of immunoglobulin in normal milk are low and depend on the degree of vascular permeability of the udder tissues (Coulon *et al.*, 2002 and Henry *et al.*, 2007). During inflammation this permeability barrier is broken, immunoglobulin concentrations increase in secretions from infected glands.

The increase in milk immunoglobulins may be effective in reducing severity of mastitis (Nickerson, 1985; and Persson, 1992). The major function of immunoglobulins is opsonization of microorganisms for phagocytosis, and they are believed to prevent bacterial adherence to epithelial membranes, inhibit multiplication and neutralize toxins. The decrease in α -lactalbumin, β -lactoglobulin associated with the sub clinical mastitis agreed with results of previous investigations reported by (Ishikawa and Shimizu, 1982). This could be due to both inflammatory damage of the mammary secretory tissues and destruction of the blood milk permeability barriers which restrict and discriminate in transfer of protein from interstitial fluid into milk.

The percentages of α casein, β casein, κ casein, and γ casein in mastitic samples were lower than those of healthy normal milk. This result may be explained by that milk from mastitic udders exhibits greatly increased proteolytic activity (Le Roux *et al.*, 1995). Plasmin is the most

important protease in milk from healthy udders but the non-plasmin proteases become more important with increasing severity of udder inflammation. This proteolysis leads to a decrease in the relative proportion of caseins (Auldism *et al.*, 1996, Weinbreck, *et al.*, 2004 and Henry *et al.*, 2007).

CONCLUSION

Although the CMT is a quick, easy to perform test and accurately identify subclinical mastitic cases it is not specific for specific pathogens.

Identification of the causative agent is an essential step in the application of antimicrobial therapy and herd management.

Infected animals should be identified as rapid as possible as it is the potential source of infection to non-infected animals.

Protein percent of milk serum whey protein, immunoglobulin, and albumin increased during subclinical mastitis. While α -casein, κ casein, and β -casein decreased in sub clinical mastitis.

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