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MICROBIOLOGICAL RESEARCH ON SOME DAIRY PRODUCTS

(With 6 Tables)

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دراسة ميكروبيولوجية على بعض منتجات الألبان

صلاح فتحي أحمد عبد العال

أجريت هذه الدراسة علي مائة عينة من منتجات الألبان (25 من كل من اللبن المعامل حرارياً بطريقة UHT والزابادي والزابادي بالفلكهة والجبن الأبيض الطري المجمعة من المحلات المختلفة بمدينة هوهنايم – شتوتجارت – ألمانيا. وتم الفحص بمعامل الجامعة لتحديد العدد الكلي للميكروبات بالإضافة إلي تواجد ميكروبات التسمم الغذائي بهذه العينات. وأسفر الفحص عن أن متوسط العدد الكلي للميكروبات في عينات اللبن المعامل حرارياً بطريقة UHT والجبن الأبيض الطري كان $4 \times 10^2,9$ و $7,8 \times 10^4$ / مللي أو جرام علي التوالي. كما تم عزل ميكروبات المكورات العنقودية والمكورات السبحية المعوية و الإيشيريشيا كولاي و كلوستريديم بيرفرنجنيز بنسبة (0,0 و 28,0 و 40,0 و 64,0 %) و (0,0 و 16,0 و 20,0 و 36,0 و 48,0 %) و (0,0 و 20,0 و 28,0 و 28,0 %) و (0,0 و 8,0 و 0,0 و 12,0 و 0,0 %) من العينات المفحوصة علي التوالي وكان متوسط عدد هذه الميكروبات كالاتي ($1,2 \times 10^3$ و $1,3 \times 10^3$ و $3,4 \times 10^3$ و 1×10^3 و $1,4 \times 10^3$ و $1,98 \times 10^3$ و $1,95 \times 10^3$ و (0,0 و $0,7,6 \times 10^2$ و $2,2 \times 10^3$ و $2,1 \times 10^3$ و $0,9 \times 10^3$ و 0,0 و $0,0 \times 10^3$) / مللي أو جرام علي التوالي وأهم المعزولات التي تم تصنيفها في هذه العينات هي ميكروب المكور العنقودي الذهبي والمكور العنقودي من نوع إبيدريميس والمكور السبحي المعوي من نوع فيكالس ومن نوع فيشيوم ومن نوع ديورانس والإيشيريشيا كولاي وكلوستريديم بيرفرنجنيز بنسبة (0,0 و 21,7 و 12,9 و 20,4 %) و (0,0 و 26,2 و 29,0 و 26,5 %) و (28,6 و 21,7 و 19,4 و 10,2 %) و (42,8 و 8,7 و 12,9 و 14,3 %) و (0,0 و 0,0 و 3,2 و 8,2 %) و (0,0 و 21,7 و 22,6 و 14,3 %) و (28,6 و 0,0 و 0,0 و 6,1 %) من إجمالي المعزولات. في حين أن كاميليو باكتري جيجينايا والسالمونيلا لم يتمكن من عزلها من أي من العينات المفحوصة. وقد تمت دراسة أهمية هذه الميكروبات الصحية وإمكانية التحكم في نقلها من الحيوان للإنسان والتسبب في الإصابة بالأمراض بالإضافة إلي الشروط الصحية الواجب توافرها لإنتاج منتج صحي عالي الجودة خالي من الأمراض.

SUMMARY

One hundred samples of dairy products (25 each) of UHT milk, plain yoghurt, fruit yoghurt and white soft cheese samples were examined for total viable count, and the presence of foodborne pathogenic microorganisms. The results declared that the mean total bacterial counts/ml or gm were 2.9×10^4 and 7.8×10^4 in examined UHT milk and white soft cheese samples, respectively. *Staphylococci*, *Enterococci*, *E. coli*, and *Clostridium perfringens* were detected in (0.0, 28.0, 40.0 and 64.0 %) & (16.0, 20.0, 36.0 and 48.0 %) & (0.0, 20.0, 28.0 and 28.0 %) and (8.0, 0.0, 0.0 and 12.0 %) of examined samples, respectively. The mean values of isolated organisms/mL or gm were (0, 1.2×10^3 , 1.3×10^3 and 3.4×10^3) & (1×10^3 , 1.4×10^3 , 1.98×10^3 and 1.95×10^3) & (0, 7.6×10^2 , 2.2×10^3 and 2.1×10^3) and (0.9×10^3 , 0, 0 and 3.3×10^3), respectively. The predominant isolated bacterial stains were *Staphylococcus aureus*, *Staphylococcus epidermidis*, *Enterococcus faecalis*, *Enterococcus faecium*, *Enterococcus durans*, *E. coli*, and *Clostridium perfringens* by frequency distribution of (0.0, 21.7, 12.9 and 20.4 %) & (0.0, 26.2, 29.0 and 26.5 %) & (28.6, 21.7, 19.4 and 10.2 %) & (42.8, 8.7, 12.9 and 14.3 %) & (0.0, 0.0, 3.2 and 8.2 %) & (0.0, 21.7, 22.6 and 14.3 %) and (28.6, 0.0, 0.0 and 6.1 %) of total isolates, respectively. Meanwhile, *Campylobacter jejuni* and *Salmonellae* failed to be detected in all examined samples. The sanitary and public health importance of these organisms as well as control measures to improve the quality of dairy products and to safeguard the consumers from infection were discussed.

Key words: *Microbiological, quality dairy products*

INTRODUCTION

Dairy products have generally been considered an excellent source of high-quality protein, calcium, potassium, phosphorus, magnesium, zinc, and the B vitamins riboflavin, niacin, vitamin B-6, and vitamin B-12 (Buttriss, 1997).

Many methods of food preservation are used for ensuring microbiological safety, among which UHT. In UHT processing, the milk is heated before its packaging and then sealed into sterilized containers in a sterile environment. The use of higher temperature 135-142°C for few seconds will increase the microbial death rate more than the loss of milk quality associated with thermal reactions (Adams and Moss, 1995). UHT seems a very promising technique as it offers numerous

opportunities for developing new stable milk with extended shelf life, high nutritional value and excellent organoleptic characteristics. UHT milk minimally processed but safe for consumers and the product requires no refrigeration (Fonberg–Broczek *et al.*, 1999).

Yoghurt is defined by the coagulated milk product that results from the fermentation of lactic acid in milk by *Lactobacillus bulgaricus* and *Streptococcus thermophilus* (Bourlioux and Pochart, 1988). Components of the human intestinal microflora and of the food entering the intestine may have harmful or beneficial effects on human health. Abundant evidence implies that specific bacterial species used for the fermentation of dairy products such as yoghurt and selected from the healthy gut microflora have powerful antipathogenic and anti-inflammatory properties. These microorganisms are therefore involved with enhanced resistance to colonization of pathogenic bacteria in the intestine, which has led to the introduction of novel modes of therapeutic and prophylactic interventions based on the consumption of monocultures and mixed cultures of beneficial live microorganisms as "probiotics" (Guarner and Schaafsma, 1998).

Cheese is an excellent food containing a wide variety of easily digested nutrients; its importance in human nutrition lies not only in supplying the consumer with a high quality protein, but also, it is rich in the essential amino acids. It contains high fat percentage, which supplies the consumer with the essential fatty acids, phospholipids and energy. Cheese is extremely an economical food, more than 90 % of the assimilated material being changed into body tissues or energy and because cheese contains all essential amino acids, it is considered the main protein supplement to farmers and most people. A great variety of cheeses are produced through traditional dairy technologies and widespread all over the world. Cheese quality is greatly influenced by the complex microbial flora from the raw milk which together with that arising from the processing environment, contributes to milk acidification, curd production and ripening often leading to final products with distinctive flavours and taste (Mauriello *et al.*, 2003). However, it can be contaminated by pathogenic m.os., thus compromising the safety of cheese and representing a hazard for consumers (Maguire *et al.*, 1991).

Milk and products derived from milk of dairy cows can harbor a variety of microorganisms and can be important sources of foodborne pathogens (Oliver *et al.*, 2005). Milk products are an excellent environment for growth of pathogenic microorganisms, which may

cause foodborne diseases. Quality and shelf life of these products depends greatly on the properties of the m.os. contaminating the product. Despite the introduction of food standards obligatory in EU countries, epidemiologists believe that 75 % of foodborne diseases are caused by bacteria (CAC, 2003). For this reason, the control of microorganisms is an important aspect of the food quality and safety.

Staphylococci, *Enterococci*, *Escherichia coli*, *Salmonellae*, *Clostridium perfringens*, and *Campylobacter jejuni* are the most important foodborne pathogens that are widely distributed throughout the environment. They have been associated with sever food poisoning outbreaks and often found in milk and dairy products (Condera *et al.*, 2004). The aim of this study was to estimate the pathogenic bacteria that contribute a major threat to the safety of dairy products and the involvement of these dairy products in foodborne diseases.

MATERIALS and METHODS

Collection of samples:

One hundred samples of dairy products (25 each) of UHT milk, plain yoghurt, fruit yoghurt and white soft cheese samples were collected in their retail packs from different localities, markets and shops in Stuttgart, Germany. The collected samples were transferred in an insulated ice-box to the laboratory with a minimum of delay for examination.

Preparation of samples:

A- UHT milks (Longeveld, *et al.*, 1976): Each sample of UHT milk was thoroughly mixed before being subjected to bacteriological examination. The surface of the retail packs was thoroughly swabbed with 70 % alcohol. A sterile, single-service hypodermic needle of syringe inserted through the package wall for bacteriological examination.

B- Yoghurt samples (A.P.H.A., 1992): Eleven (11) grams of the prepared sample were added to 99 ml of sterile distilled water (40-45°C), shaken until a homogenous dispersion is obtained to obtain dilution 1/10 from which decimal dilutions were prepared using a sterile peptone water solution 0.1 %.

C- Cheese samples: (A.P.H.A., 1992): Eleven (11) grams of cheese sample was transferred aseptically to a sterile blender then add 99 ml sterile, freshly prepared aqueous solution of 2 % Na citrate (sodium citrate) at 40-45°C. Mix for 2 min and invert the blender container to rinse particles from the interior wall then remix for approximately 10

sec. Mix thoroughly till complete emulsification to make dilution 1:10, from which decimal dilutions were prepared using a sterile peptone water solution 0.1 %.

Microbiological examination: The prepared samples were subjected to the following examination:

1- Preparation of serial dilution (A.P.H.A., 1992): One ml of each prepared sample was added to 9 ml of sterile peptone water solution 0.1 % to make serial decimal dilutions.

2- Total colony count (A.P.H.A., 1992): One ml from the previously prepared dilution was inoculated onto duplicates of standard plate count (SPC) agar and incubation at 37°C for 24 h.

3- Staphylococci count (Chapman, 1945): 0.1 ml from the previously prepared dilutions of the examined samples was transferred and evenly spread on the surface of Mannitol salt agar medium (Oxoid, 1990) plates. Inoculated plates were incubated at 37°C for 48 h. and Staphylococci count/mL was calculated and recorded.

4- Enterococci count (Gelsomino *et al.*, 2003): 0.1 ml from the previously prepared dilutions of the examined samples was inoculated on the surface of kanamycin esculin azide agar (kAA; Merck, Darmstadt, Germany) at 37°C for 24 h.

5- *E. coli* count (Bacteriological Laboratory of Hygiene and Environment Institute): 0.1 ml from the previously prepared dilutions of the examined samples was spread onto Targitol medium and Endo medium (Oxoid, 1990) then incubation at 37°C for 24 h.

6- Isolation of Salmonella spp. (Jayarao and Henning, 2001):

*Pre-enrichment: 25 ml from the previously prepared sample added to 225 ml peptone water with Novobiocin (Standard: 100 mg/1 ml sterile D.W. & Test: 0.9 ml prepared solution/225) then incubated at 37°C for 24 h.

*Selective enrichment: One ml peptone water from previously prepared pre-enrichment added to 10 ml Rappaport Vassiliadis broth, Difco Laboratories (Two tubes) after that one tube incubated at 37°C for 24 h. and the second at 43°C for 24 h.

*Plating on selective medium: 0.1 ml of incubated Rappaport evenly spread on the surface of Xylose Lysine Desoxycholate (XLD) and Brilliant Green Phenol Red Lactose Sucrose (BPLS) plates, Unipath Co. Inoculated plates were incubated at 37°C for 24 h. and 43°C for 24 h.

7- Anaerobic spore formers count (Bacteriological Laboratory of Hygiene and Environment Institute): Using Thioglucolate medium (Oxoid, 1990).

*1 ml of prepared sample in 3 tubes of Thiogluconate, heating at 70°C for 20 min and incubated at 37°C for 24 h. anaerobically (anaerobic jar with anaerobic anaerocult A sachet moisted with 17 ml D.W.).

*1 ml from the previously prepared serial dilution in 3 tubes of Thiogluconate, heating at 70°C for 20 min and incubated at 37°C for 24 h. anaerobically.

*Loopfuls from previously incubated tubes were streaked on blood glucose agar plate. The inoculated plates were incubated at 37°C for 48 h. anaerobically. Specific colonies grew surrounded by large zone of haemolysis.

8- Isolation and Identification of Campylobacter (Hunt *et al.*, 2001):

*Selective enrichment: 1 ml from prepared sample and 9 ml Preston selective enrichment broth. Then incubated at 43°C for 48 h. in microaerophilic atmosphere (anaerobic jar with anaerocult C sachet of microaerophilic organism moisted with 6 mL D.W.), Oxoid Ltd., Basing Stoke, UK.

*Plating on selective medium: 0.1 ml of Preston broth enrichment put on filter type AC(pore size, 0.45 µm) on the surface of Campylobacter agar medium (Columbia Agar Base + Horse Blood + Campylobacter Selective Supplement Cod SR 204 E + Campylobacter Growth Supplement Code SR 084 E) incubation at 37°C for 2 h. then remove the filter. Incubation of the media at 43°C for 48 h. microaerophilic in anaerobic jar.

Identification of isolated organisms: Purified colonies were identified by using colony morphology, gram staining characteristics, oxidase, Catalase, coagulase production and biochemical reactions. Specific identifications were made using Commercial micro methods (API Staph for Staphylococci, API 20 Strept for Enterococci, API 20 E for E. coli, API 20 A for Clostridia and API Campy, Bio Merieux, France). Specific Serological tests were made for Salmonellae spp.: Polyvalent (I or II) and Monovalent.

RESULTS

Table 1: Statistical analytical results of total bacterial counts/ml in examined UHT milk and White soft cheese samples.

Examined samples	Total No. of samples	Positive samples		Min.	Max.	Mean	±S.E.M.
		No.	%				
UHT milk	25	10	40.0	5×10^3	5×10^4	2.9×10^4	0.36×10^4
White soft cheese	25	18	72.0	2×10^4	23×10^4	7.8×10^4	1.1×10^4

* -ve samples > 100

Table 2: Statistical analytical results of Staphylococci counts/ml in examined dairy product samples.

Examined samples	Total No. of samples	Positive samples		Min.	Max.	Mean	±S.E.M.
		No.	%				
UHT milk	25	0	0.0	0	0	0	0
Plain yoghurt	25	7	28.0	4×10^2	2×10^3	1.2×10^3	0.11×10^3
Fruit yoghurt	25	10	40.0	6×10^2	3.3×10^3	1.3×10^3	0.17×10^3
White soft cheese	25	16	64.0	8×10^2	9×10^3	3.4×10^3	0.41×10^3

* -ve samples > 100

Table 3: Statistical analytical results of Enterococci counts/ml in examined dairy product samples.

Examined samples	Total No. of samples	Positive samples		Min.	Max.	Mean	±S.E.M.
		No.	%				
UHT milk	25	4	16.0	2×10^2	2.3×10^3	1×10^3	0.19×10^3
Plain yoghurt	25	5	20.0	5×10^2	2.6×10^3	1.4×10^3	0.19×10^3
Fruit yoghurt	25	9	36.0	8×10^2	3.5×10^3	1.98×10^3	0.15×10^3
White soft cheese	25	12	48.0	1.2×10^3	4.9×10^3	1.95×10^3	0.20×10^3

* -ve samples > 100

Table 4: Statistical analytical results of E. coli counts/ml in examined dairy product samples.

Examined samples	Total No. of samples	Positive samples		Min.	Max.	Mean	±S.E.M.
		No.	%				
UHT milk	25	0	0.0	0	0	0	0
Plain yoghurt	25	5	20.0	3 x10 ²	13 x10 ²	7.6 x10 ²	0.74 x10 ²
Fruit yoghurt	25	7	28.0	12 x10 ²	3.3 x10 ³	2.2 x10 ³	0.15 x10 ³
White soft cheese	25	7	28.0	14 x10 ²	5.2 x10 ³	2.1 x10 ³	0.27 x10 ³

* -ve samples > 100

Table 5: Statistical analytical results Clostridium counts/ml in examined dairy product samples.

Examined samples	Total No. of samples	Positive samples		Min.	Max.	Mean	±S.E.M.
		No.	%				
UHT milk	25	2	8.0	1 x10 ²	1.7 x10 ³	0.9 x10 ³	0.22 x10 ³
Plain yoghurt	25	0	0.0	0	0	0	0
Fruit yoghurt	25	0	0.0	0	0	0	0
White soft cheese	25	3	12.0	2.5 x10 ³	4 x10 ³	3.3 x10 ³	0.15 x10 ³

* -ve samples > 10

Table 6: Frequency distribution of isolated bacterial stains in examined dairy product samples.

Isolated bacterial stains	UHT milk (n=25)		Plain yoghurt (n=25)		Fruit yoghurt (n=25)		White soft cheese (n=25)	
	No. of isolates	F.	No. of isolates	F.	No. of isolates	F.	No. of isolates	F.
<i>Staph. Aureus</i>	0	0.0	5	21.7	4	12.9	10	20.4
<i>Staph. Epidermidis</i>	0	0.0	6	26.2	9	29.0	13	26.5
<i>Entrococcus faecalis</i>	2	28.6	5	21.7	6	19.4	5	10.2
<i>Entrococcus faecium</i>	3	42.8	2	8.7	4	12.9	7	14.3
<i>Entrococcus durans</i>	0	0.0	0	0.0	1	3.2	4	8.2
<i>E. coli</i>	0	0.0	5	21.7	7	22.6	7	14.3
<i>Clostridium perfringens</i>	2	28.6	0	0.0	0	0.0	3	6.1
<i>Campylobacter jejuni</i>	0	0.0	0	0.0	0	0.0	0	0.0
<i>Salmonellae</i>	0	0.0	0	0.0	0	0.0	0	0.0
Total	7	100.0	23	100.0	31	100.0	49	100.0

DISCUSSION

Results presented in Table 1 showed that the total bacterial count could be detected in examined UHT milk and white soft cheese samples at varying percentages 40.0 and 72.0 %, with a mean counts/ mL or gm of 2.9×10^4 and 7.8×10^4 , respectively. Lower prevalence in UHT milk was reported by Schaal and Noecker, 1977.

It is evident from the previously mentioned data that UHT milk samples yielded viable bacteria in 10 out of 25 samples and this normally unacceptable because sterilization or UHT treatment of milk are essential to ensure total microbial safety and enzymatic stability of milk (Korhonen *et al.*, 1998) and allow prolonged self life of milk up to 6 months but the quality of packing especially its permeability proves to be very important (Reddy and Love, 1999). Also, mesophilic aerobic spore formers have ability to produce highly heat resistant spores that may survive sterilization or UHT treatment (Pettersson *et al.*, 1996).

The higher total count was observed in examined white soft cheese samples due to several factors as the extent of handling in its production (Seligman, 1976), post pasteurization contamination, usually through contact with equipment surfaces or from the air and biofilms residing on surfaces which are one potential source of contamination (Austin and Bergeron, 1995).

Table 2 revealed that *Staphylococci* failed to be detected in UHT milk samples and could be isolated from 28.0, 40.0 and 64.0 % from the examined plain yoghurt, fruit yoghurt and white soft cheese samples with a mean values/ gm of 1.2×10^3 , 1.3×10^3 and 3.4×10^3 , respectively. The predominant isolated *Staphylococci* stains were *Staphylococcus aureus* and *Staphylococcus epidermidis* by frequency distribution of (21.7, 12.9 and 20.4 %) & (26.2, 29.0 and 26.5 %) of total isolates, respectively (Table, 6). Lower prevalence of *Staphylococci* in yoghurt was obtained by Arnott *et al.*, 1974 while, in cheese lower incidences were recorded by Bastepe and Kösker, 1981; Papageorogiou *et al.*, 1998 and Dipietro *et al.*, 2004. Nearly similar count of *Staphylococci* in cheese was obtained by Ashenafi, 1990 while, higher counts were recorded by Mol and Vincentie, 1975 and Turantas *et al.*, 1989 and a lower count was detected by Bastepe and Kösker, 1981.

The absence of *Staphylococci* in UHT milk samples due to the fact that they are heat labile microorganisms so destructed by heat but the biggest problem is the toxins which are heat stable so in UHT milk it is advisable to search about toxin of *Staphylococci* rather than the organism (Dunstall *et al.*, 2005). The low prevalence of *Staphylococci* in yoghurt samples explained by the antagonistic effect of yoghurt against

representative of conditionally pathogenic intestinal microorganisms as *Proteus*, *Klebsiella*, *Para coli* and *Staph. aureus* (Kockova *et al.*, 1980).

Results summarized in Table 3 declared that the *Enterococci* could be detected in 16.0, 20.0, 36.0 and 48.0 % of examined UHT milk, plain yoghurt, fruit yoghurt and white soft cheese samples, respectively. The mean counts/mL or gm of *Enterococci* in examined samples were 1×10^3 , 1.4×10^3 , 1.98×10^3 and 1.95×10^3 , respectively. The main isolated *Enterococci* stains were *Enterococcus faecalis*, *Enterococcus faecium* and *Enterococcus durans* by frequency distribution of (28.6, 21.7, 19.4 and 10.2 %) & (42.8, 8.7, 12.9 and 14.3 %) and (0.0, 0.0, 3.2 and 8.2 %) of total isolates, respectively (Table, 6). The results of UHT milk samples in agreement with that reported by Schaal and Noecker, 1977 who supposed that 10 % of UHT milk packages proved to be faulty. Nearly similar percentage was detected in yoghurt samples by Arnott *et al.*, 1974. While, in cheese samples Gelsomino *et al.*, 2002 and Huys *et al.*, 2003 could isolate the same *Enterococci* strains. Mikulec and Jovanovic, 2005 reported lower *Enterococci* counts and higher count obtained by Ashenafi, 1990.

Enterococci can be used as indicators of faecal contamination, they have been implicated in outbreaks of foodborne illness and they have been ascribed a beneficial or detrimental role in foods. *Enterococci* may survive heat processing and cause spoilage of the products as well as causing food infection (Franz *et al.*, 1999).

Results tabulated in Table 4 revealed that *E. coli* failed to isolated from examined UHT milk samples. While, it could be isolated from 20.0, 28.0 and 28.0 % of the examined plain yoghurt, fruit yoghurt and white soft cheese samples with a mean counts/ gm of 7.6×10^5 , 2.2×10^3 and 2.1×10^3 , respectively. The *E. coli* isolated stains represented 21.7, 22.6 and 14.3 % of total isolates, respectively (Table, 6). Arnott *et al.*, 1974 reported lower percentage of *E. coli* in yoghurt samples. Meanwhile, Dipietro *et al.*, 2004 and Brenda *et al.*, 2005 obtained lower prevalences of *E. coli* in cheese and higher results were recorded by Ashenafi, 1990 and Papageoriou *et al.*, 1998.

E. coli failed to be detected in UHT milk samples because the main source of this microorganism appear to be the intestinal tract of cattle (Bilici and Tayfur, 2002) and can't survive efficient sterilization or UHT treatment (Pettersson *et al.*, 1996). The lower percentage of *E. coli* in yoghurt is due to acid adaptation (pH 3.6-3.9) which reduce survival of *E. coli* (Yi and Chou, 2001).

Table 5 showed that *Clostridium perfringens* failed to isolated from examined yoghurt samples. While, could be isolated from 8.0 and 12.0 % of examined UHT milk and white soft cheese samples with a mean counts/ml or gm of 0.9×10^3 and 3.3×10^3 , respectively. *Clostridium perfringens* isolated stains represented 28.6 and 6.1 % of total isolates, respectively (Table, 6). Turantas *et al.*, 1989 could not isolate *Clostridium perfringens* from cheese while, Papageoriou *et al.*, 1998 could isolate *Clostridium perfringens* with high percentages.

UHT milk samples contain *Clostridium perfringens* and this related to *Clostridium* is anaerobic spore formers so when spores present with faulty storage it will be grow and produce serious defects as putrefaction and acidification or other anomalies of taste (Schaal and Noecker, 1977). *Clostridium perfringens* could not detected in yoghurt and this may be due to increase lactic acid and titrimetric acidity as well as by a decrease in pH value (Pazakova *et al.*, 1997).

Results mentioned in Table 6 reported that *Campylobacter jejuni* failed to be detected in the examined samples. The spreading of *Campylobacter jejuni* is mainly associated with food products of animal origin but products were always negative as a result of better controls in the processing of these products (Baffone *et al.*, 1995).

Also, it is observed from this Table that *Salmonellae* failed to be detected in all examined samples. Nearly similar findings were obtained by Turantas *et al.*, 1989 and Papageoriou *et al.*, 1998 while, Grieger *et al.*, 1976 found *Salmonellae* in milk products mainly cheese and explained this as a result of rennet produced by one enterprise. Also, Colak *et al.*, 2007 could isolate *Salmonellae* from cheese.

Contamination of milk and dairy products by pathogenic microorganisms can be endogenous origin, following excretion from the udder of an infected animals or may be also of exogenous origin, through direct contact with infected herd or through environment (water & personnel). Heat treatment and processing of milk can inhibit or encourage the multiplication of microorganisms. Deficiencies in the hygienic measures of milk and dairy products storage, particularly refrigeration and in the HACCP plan that was not properly implemented should be corrected. It is important to inspect the manufacturing plant than to examine the single dairy product on the market.

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