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Fungal Contamination of White Cheese at the Stage of Consumption in Saudi Arabia

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Abstract: Thirteen species appertaining to 4 genera of molds and yeast were isolated from twenty cheese samples using malt extract and 10% NaCl-Czapek's gar media at 25°C. The results revealed that the white cheese under examination were less polluted and most of the samples tested, were free of fungi comparing with other findings. The total fungal counts in cheese samples were relatively low and ranged from 95 and 125 colonies / 160 cheese pieces on the two isolation media, respectively. Members of Penicillium (P. brevicompactum, P. chrysogenum, P. paxilli and P. waksmanii) and Aspergillus (A. flavus, A. niger and A. oryzae) were the most prevalent fungi on cheese samples. Also, some unknown yeasts were encountered, in moderate occurrence. The preliminary examination, using 70 isolates, was done to test their ability to produce protease enzyme. The results indicated that 59 fungal isolates showed their ability to hydrolyze milk protein. Among the positive isolates 24 could produce the enzyme in high amount, while 20 produce moderate, and remaining 15 produce weak amounts.

Key words: Cheese, fungal contamination, Aspergillus, Penicillium

Introduction

Cheese is considered as one of the most important foodstuffs consumed by human and it contains a source of high quality animal protein having all the essential amino acids (El-Shrief, 2000). It is also a rich source of calcium, phosphorus and many other micronutrients. Cheese can easily become molded by a large number of species of fungi and a group of unicellular fungal organisms (yeast) during ripening, after cutting and slicing and during storage in shops or at home (Carter & Cole, 1990). Fungi are undesirable organisms which can grow, at a wide range, causing spoilage as well as are responsible for flavour defects, discoloration and poor appearance of the product (Walker, 1977). Furthermore, they produce toxic metabolites (mycotoxins) which may penetrate the cheese and polishing its surface might not sufficiently remove them and effect the health of the consumers (Abdel Sater et al., 1995; Lund et al., 1995; El-Sharief, 2000).

Proteases are one of the most important groups of enzymes constituting two thirds of the total industrial enzymes marketed (Gehartz, 1990; Singh et al., 1994). They play an important role in the food processing industry (Wu & Hong, 1998). These enzymes have been isolated and characterized from mammals, plants, bacteria, yeasts and fungi (Nielsen, 1991; Beuchat & Pitt, 1992).

The main objective of this work was to establish the distribution of fungi in different types of cheese commonly consumed in Saudi Arabia. Also, evaluation of proteolytic activity by several strains of fungi was detected.

Materials and Methods

Twenty different cheese samples were collected from various markets in Riyadh-City, Saudi Arabia, (Table 1). The samples were transferred to the laboratory and kept in a refrigerator (6-8°C) till fungal analysis.

Estimation of fungi: The plating technique method was employed for this experiment. Four pieces (about 0.5 g) from each sample were scattered on the surface of plates containing solidified malt extract agar (g/L: malt extract, 20; peptone, 5; agar, 15) and 10% NaCl-Czapek's agar media (g/L; glucose, 10; NaNO₃ 3; KH₂PO₄. 7H₂O; MgSO₄, 0.5; KCl, 0.5; agar, 15). Four plates for each sample (2 for each medium) were used. The plates were incubated at 25°C for 1-2 weaks and the developing fungi were counted, identified and calculated per 8 pieces for each sample.

Table 1: Number of sample and kind of cheese under examination

Tubic 1.1	tamber of sample and kind of cheese ander examination
No.	Type of cheese
1	Pinar
2	Kraft light (Phildelphia)
3	Danish feta (Danland)
4	La Vache qui rit (cream)
5	Danish feta (Almarai)
6	Kraft (Milky Cheese spread)
7	Franch feta (Valbreso)
8	Danish white (The three cows)
9	La vache qui rit (Savoury)
10	Strong covv
11	Cream cheese (Almarai)
12	Kraft (cream Spread)
13	Mozzarella
14	Al Fallaha Labneh
15	Danish (Mozzarella
16	Egyptian cheese
17	Saudi Arabian Cheese
18	Syrian haloom Cheese
19	Turkish cheese
20	Czach chasca

Proteolytic activity: The prevalent species, in the current study, were screened for their ability to produce protease enzyme using the procedure of El-Gendy (1966). The medium of the following composition (g/L) was used. Peptone 10, agar 20, pH 6.8. After autoclaving the medium, sterile skim milk (10% solution of powder deffated milk in water was sterilized separately) was added at the rate of 5 ml per 100 ml medium before pouring in plates. Test fungi were inoculated onto the center of the plates and incubated at 28°C for 10 days. Complete degradation of milk protein was seen as clear zone around the colonies. The clear zones were recorded.

Results and Discussion

The mycological analysis of the most consumed white cheese in Saudi Arabia, revealed that the samples were less contaminated and most of the samples tested were free from fungi comparing with other findings (Table 2). Thirteen species representing 4 genera of moulds and yeast were isolated from the twenty cheese samples tested using platebating technique on malt extract (8 species + 3 genera) and 10% NaCl-Czapek's agar (12 and 4) at 25°C. The fungal total counts in cheese samples were relatively low and ranged from 95 to 125 colonies/160 cheese pieces on the two isolation media, respectively. The lowest incidence of fungal propagules in tested cheese signifying the accurate methods of processing, pickling and preservation techniques of mainly cheeses. Also, due to continuous sampling of different food

products to be analyzed for microbiology and get rid of positive lots and the severe restrictions and rigorous laws made by Saudi government on export food and foodstuffs. The present results are in contrast with other reports on milk and milk products in many parts of the world. For instance, the counts of moulds in Swiss cheese, as reported by Bullerman (1976), ranged from 10-750 colonies/g. Kivance (1990) examined 25 random samples of fresh Van herby and pickled white Turkish cheeses for moulds and aflatoxins. He observed that the mean values of the total moulds were 2.5x105 and 4.95x104 colonies/g, respectively. Also, the same author (1992) tested hard cheese (surface and core) for fungal contamination and found that the mean total mould counts were 3.02x10¹⁰ and 3.02x10³ colonies/g, respectively. Lund et al. (1995) identified 371 fungal isolates from hard, semi hard and semi soft cheeses from Denmark, France and other countries. But, in spainian cheese, Arizcum et al. (1996) found that all mould counts were < 1000 colonies/g and they could identify 9 genera from the samples examined. In Egyptian cheese, El-Bassiony et al. (1980) isolated 11 genera of moulds. El-Sawi et al. (1994) examined buffaloes milk cottage and isolated 35 species belonging to 9 genera. Also, in (1995) Abdel-Sater et al. and Saad & Hemida isolated respectively 26 and 29 species belonging to 16 and 8 general from four types of local Egyptian and roqueforti cheeses. The previous authors indicated that these cheeses were heavily polluted with micro-organisms with counts of 3500-11000 colonies/g cheese. More recently, El-Shrief, in Egypt (2000) examined 260 samples representing 5 kinds of cheese locally manufactured.

The results in Table 2 indicated that members of Penicillium and Aspergillus were the mainly contaminated moulds of all the tested samples. These results were greatly identical to those obtained by Polonelli et al. (1984), Pitt & Hocking (1985), Scott (1989), Abdel-Sater et al. (1995), Taniwaki & Dender (1992); Arizcum et al. (1996), El-Shrief (2000). They indicated that the majority of moulds isolated from cheeses and other foodstuffs consisted of Penicillium and Aspergillus. Penicillium was the first common genus in tested samples. It was occurred in 56 and 59% of the samples contributing 56.8 and 66.7% of total fungi on malt extract and 10% NaCl-Czapek's agar, respectively. From this genus, 7 species were identified of which P. brevicompactum, P. chrysogenum, P. paxillii and P. waksmanii were the most prevalent. They were present in 25-45% and 20-55% of the samples on the two isolation media. The remaining three Penicillium species were isolated in rare occurrence at least on one medium and not on the other (P. corylophillum and P. frequentans on 10% NaCl, but P. roquefortii on malt extract) at 25°C (Table 2). The above species and other Penicillium species were also, isolated, with variable incidences, from different cheese samples all over the world (El-Malt, 1993; Hassanin, 1993; El-Sawi et al., 1994; Abdel-Sater et al., 1995; Lund et al., 1995, Larsen, 1997; El-Sharief, 2000).

Asperaillus (5 species) was the second dominant genus. emerging in 53% and 58% of the samples comprising 41.1 and 34.4% of total fungal isolates on malt extract and 10% NaCl-Czapek's agar, respectively. Among the 5 species isolated from the genus, A. flavus and A. niger were the most common. They were isolated from 50 and 53 and 46 and 50% of the samples on the two isolation media, respectively. A. oryzae, A. sydowii and A. terreus were less frequent and isolated each from 15 or 20% of the samples (Table 2). In this respect, several species of Aspergillus were isolated from raw milk or milk products in many parts of the world. Bullerman (1980) isolated A. flavus and A. ochraceus from domestic and imported retail samples of cheese. Aran & Eke (1987) could isolate A. versicolor and Eurotium species from Turkish Kasar cheese. Hassanin (1993) could isolate A. flavus and A. niger from 16 samples of cheese collected randomly

from several locations in Cairo and Giza, Egypt. Lund et al. (1995) isolated A. versicolor from hard, semi-hard and semi-soft cheese from Denmark, France and other countries. Also, Saad & Hemida (1995) examined 40 samples of Roquefort cheese and could isolate A. flavus and A. niger. Barrios et al. (1997) examined 52 commercial cheese produced in southern Spain and could detect A. glaucus, A. niger, A. terreus and A. flavus in most of the examined samples. Abdel-Sater et al. (1995) and could isolate A. flavus, A. niger, A. sydowii, A. tamarii, A. terreus, A. ochraceus and A. scolerotiorum from the Egyptian cheese. More recently, El-Sharief (2000) examined cheese samples and found A. flavus, A. flavipes, A. niger and A. terreus among the microorganisms contaminating the samples.

Cladosporium cladosporioides was isolated only on 10% NaCl-Czapek's agar, occurring in 15% of the samples. This specie and other Cladosporium species were very common with a world-wide distribution and occurring on foodstuffs (including cheese), soil, air, textiles, seeds, plant materials, stored fruits, cereal grains and other organic matter (Samson et al., 1995). Also, Cladosporium species were isolated from different types of cheese such as kariech (El-Bassiony et al., 1977); hard (Abdel-Rahman & El-Bassiony 1984); Teleme (Zerfiridis, 1985); Spanish (Diaz et al., 1995); Soft

Table 2: Total counts (TC per 160 pieces in all samples) and percent frequency (F%) of mycoflora isolated from examined cheese on malt extract and 10% NaCl-Czapek's agar at 25°C

Organisms	Malt extract		10% NaCl	
	TC	F%	TC	F%
Aspergillus	39	53	43	58
A. flavus Link	14	50	10	53
A. niger Van Tieghem	21	46	19	50
A. oryzae (hlb.) Cohn.	-	-	5	15
A. sydowii(Bain.&Sart.)Thom	4	15	5	20
& Church				
A. terreus Thom	-	-	4	15
Cladosporium cladosporioides				
(Fres.) de Vries	-	-	2	5
Penicillium	54	56	80	59
P. brevicompactum Dierckx	7	30	10	20
P. chrysogenum Thom	18	45	17	35
P. corylophilum Dierckx	-	-	3	5
P. frequentans Westing	-	-	2	5
P. paxilli Bain.	3	25	11	25
P. roguefortiiThom	14	20	-	-
P. waksmanii Zaleski	12	40	37	55
Yeasts	11	30	15	45
Total counts	95		125	
Number of genera = 4	3		4	
Number of speices = 13	8		12	

(Sympayo et al., 1995) and Egyptian cheeses (El-Sharief, 2000). The results recorded in Table 2 show that the total counts of yeasts fluctuated between 11-15 colonies/160 pieces in all samples. They occurred in 30 and 45% of the samples contributing 11.6 and 12.0% of total fungal isolates. Yeasts were also, isolated from food or food products as indicated by several workers in many parts of the world (Mercado & Rivas, 1986; Guven et al., 1995; Abdel-Sater et al., 1994; Barakat & Abdel-Sater, 1999; El-Sharief, 2000). Testing the ability of different fungal isolates, recovered in current study, to produce protease enzymes indicated, that the most fungal isolates (59 out of 70) were able to produce these enzymes (Table 3). It was observed that not only the species of a single genus differed in the production of enzymes but also, the different isolates within the same species. Of 70 isolates tested 48.3 % could be realized as producers of protease enzymes. Among the positive isolates, 24 (40.7%) belonging to Aspergillus niger, Penicillium roquefortii and P. waksmanii achieved high activity and

Table 3: Protease	production	by common	fungal	isolates tested.
Organisms	NIT	NIP	Degree	of production

Organisms	1311	1.411	Dog	Begiec of production	
			+	+ +	+ + +
Aspergillus flavus	10	10	1	2	7
A. niger	8	8	2	3	3
A. sydowii	9	7	3	2	2
Penicillium	6	5	1	3	1
brevicombactum					
P. chrysogenum	11	9	2	4	3
P. paxilli	4	2	1	1	-
P. roguefortii	15	12	3	4	5
P. waksmanii	7	6	2	1	3
Total isolates	70	59	15	20	24

NIT = number of isolates tested; NIP = number of isolates positive; +: weak producers; ++: moderate producers;

+++: high producers.

these . Isolates of P. brevicompactum and P. chrysogenum proved to be moderate producers whereas, isolates apperataining to A. sydowii and P. paxilli could produce the enzyme with weak amounts (Table 3). These results were in agreement with those of Moharram and El- Zavat (1989). They noticed that most of isolates recovered from Tilapia fish scales could produce the protease enzymes. More recently Omar et al. (1999) reported 72 out of 96 isolates to be proteolytic including 28 isolates highly proteolytic, 21 moderately degrading milk protein and the remaining isolates were non proteolytic. Also, protease enzymes were isolated and characterized from differrent fungal species as reported by several workers (Desphpande, 1992; Singh et al., 1994; Simpanya and Baxter, 1996; Staszczak et al., 1996; Lopez -Diaz et al., 1996; Wu and Hang, 1998).

In conclusion, most cheese samples were free from fungal species, since the government makes severe restrictions and rigorous laws on foodstuff products. Also, good processing and preservation techniques of different products lessen to great extent any contamination with microorganisms. For these and other reasons cheese samples examined in the present study were less conaminated with fungi.

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