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Studies on Mycosis and Mycotoxicosis in Cattle

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ABSTRACT

The present study was conducted on forty diseased and apparently healthy cases of cattle (20 of each) at Cairo Governorate. From districts of diseased cases, 40 serum samples of these cattle and 100 samples of milk of mastitic animals, nasal swab, vaginal swabs and feeds samples which used in breeding of diseased animals (25 0f each) were collected. The mycological examination of collected samples revealed isolation of Aspergillus species in predominant rates from the above samples of infected animals, where in cattle recovered from (88%, 96%, 80%, and 100%) respectively. Whereas, other genera and species of moulds and yeast were recovered in a comparatively lower significant rates. Sera of cases of diseased cattle contained significant levels of aflatoxins and zearalenon. Meanwhile, 60% of cattle had the mean levels of aflatoxins (15.20±0.4 ppb) and zearalenon in 80% in cattle with the mean level of (62 ± 0.3). When, the used feed samples in breeding of these animals were had the amounts of AFB1; OA; ZEAR; T2; and FB1 were detected in (60%; 40%; 32%; and 44%) of feed samples, with the mean levels of (55.0±1.50, 45.0±0.30 and 31.0±0.2 and 40.0±2.1 ppb), respectively. The biochemical study on sera of the infected and those fed treated toxic feed with ammonium hydroxide 3%, demonstrated that the AFB1 and zearalenon in serum of cattle lead to a marked elevation in the serum enzymes activities of (AST, ALT and GGT, which is indicate of hepatocellular damage. The levels of serum urea and creatinine were significantly high in toxicated animals as compared to apparently healthy animals. Mycotoxins in cattle inducing significantly decreased values in serum total protein, albumin, alpha globulin, beta globulin, and gamma globulin. The mycotoxicated animals had significant increase in serum level of malondialdehyde (MDA) and serum NO level which caused induction of inducible nitric oxide synthase (iNOS) protein, resulting in cytotoxicity and DNA damage. The feeding of treated toxic feed with ammonium hydroxid resulted a significant improvement of al previous altered biochemical parameters which resulted from toxicosis. The significance of the present result was fully discussed.

Key word: Aflatoxins, zear., T2, Aspergillus sp., Biochemical parameters, Electrophoresis patterns, NO., MD.

INTRODUCTION

Up to date the environmental pollution is considered the essential cause of animal diseases particularly pollution with fungi and their toxins of the used feed and water as well as contamination of human food. Hence, great attention has been paid to the increased importance of fungi and their mycotoxins, which are serious fungal metabolites. This is in turn will contribute to the production of high quality food material for human consumption and the profitability of agricultural industry. Mycotoxins are formed by certain fungal species, whenever environmental factors are conductive during the growth of these frequently occurring mycomycetes on food stuffs and animal feeds; the process takes place as a secondary metabolism. These natural toxins have a broad spectrum effects [1-6]. Unlike many bacterial toxins, they are in general highly stable in face of environmental influences. This also means that, when foodstuffs for human consumption are preserved or prepared by the conventional heating process, the mycotoxins are capable of surviving undamaged thus reaching the eventual consumer. These fungi and mycotoxins have serious effects upon the growth rate and health of human being and animals, as some mycotoxins had been found to be carcinogenic, tremorgenic, haemorrhagic and dermatitis[7,-9]. The mycotoxins of greatest agricultural and public health significance include aflatoxins, ochratoxins, trichothecenes, fumonisins, zearalenone, and ergot alkaloids [10-12]. However, the fungi and their toxins are widely distributed through the world where they occur in soil, on plants, plants debris and similar organic subtracts. They cause significant economic losses in agriculture, morbidity and mortality in animals and immunological compromised humans, where it is capable of killing cells by causing extensive damage to cellular membrane [13,14,5]. On the other hand, epidemiological studies associated with mycotoxins had a wide range of biological effects, including pulmonary oedema in pigs and ruminants (15Harrison et al., 1990), nephrotoxicity and liver cancer in different animal species [16, 3,6, 9]. The International Agency for Research on Cancer has declared that mycotoxins are potentially carcinogenic to human. Aflatoxin B1 had been statistically associated with a high

incidence of liver cancer in certain areas of Transkei, South Africa; in China and Egypt [17, 6]. All the previous literatures recorded that the pollution affect upon the growth rate and health of human being and animals including aneamia, stunted growth, carcinogenic, tremorgenic, haemorrhagic, dermatitic, pulmonary edema, immunosuppressive and hormonal effects [3,5,6, 18]. The aim of the present work was to investigate the problem of fungal and mycotoxicosis in cattle and detection the role of feed pollution as main environmental factors of animal diseases.

MATERIAL AND METHODS

Material

Serum and feed samples:

Forty diseased cases of cattle in farms at Cairo Governorate were investigated . From districts of diseased cases,40 serum samples of cattle and 100 samples of milk of mastitic animals, nasal swab, vaginal swabs and feeds samples) which used in breeding of diseased animals(25 0f each) were collected.

Mycotoxins standards

Standerds and immunoaffinity column of AFB₁, B₂, G₁ and G_{2 and} OA, ZER, T-2 AND FB1 were purchased from Sigma Chemical Company (USA).

Mycological examination of samples

The collected samples of feeds, milk and swabs were subjected for isolation and identification of fungi as recommended by Conner et al. [19].

Detection of mycotoxins in feed and sera of diseased cattle :

Detection of mycotoxins in serum of cattle and feed stuffs by fluerometric methods as described by Hansen [20] using immune-affinity column method.

Detoxification of aflatoxicated feeds by addition of ammonium hydroxid 3%: The feed samples that contaminated over permissible limits(over 700-1000 ppm/ ten of feed) were treated by addition of ammonium hydroxid 3%. After 7 dayes of treatment, the treated feeds were given to group of apparently healthy cattle calves in comparison to other group given toxicated feeds without any treatment. The feeding was continues for I month. All the animals were tested biochemically every 10 dayes during treatment for evaluation the influence of ammonium hydroxid on the toxicities of aflatoxins on the function of vital internal organs and patterns of serum protein electrophoresis.

Biochemical investigations of sera of diseased cattle : Blood samples were collected in small labeled dry and clean vials without anticoagulant in centrifuge tube, allowed to clot and then centrifuged at 3000 rpm for 15 minutes for separation of serum which used to assay the biochemical parameters as aspartate aminotransferase (AST) and alanine aminotransferase (ALT) activities according to Reitman and Frankel, [21], serum urea according to Wybenga et al. [22], serum creatinine level according to Henry (231974), Estimation of serum total protein and electrophoretic pattern were carried out after SonnenWirth and Jaret [24] and Davis [25], respectively. Serum lipid peroxide levels of malondialdehyde (MDA) were determined according to the method described by Yoshioka et al. [26]. Blood GSH concentration was measured using the method described by Buetler et al. [27]. Serum NO levels of the cattle were measured by enzymatic Greiss reaction [28]. Estimations of vitamin E, A and C were performed according to Henry [23].

Statistical Analysis

The obtained date were computerized and analyzed for significance, Calculation of standard error and variance according to [29].

RESULTS AND DISCUSSION

The past three decades have witnessed dramatic change in man's and animal's environment and his immune defenses. Consequently, the fungi have assumed as a major role in respiratory and systemic infectious diseases. The most common isolated fungi from mastitic milk samples of cattle were Aspegillus sp. (88%), Fusarium sp. (40%), Pencilium sp.(8%), Cladosporium sp.(8%), Alternaria sp. (4%), *C.albicans* (60%), *C. tropicalis* (24%) and *Rhodotorula rubra*(16%). The percentage of moulds in raw milk samples varies depending on geographical location and season of the year. The specific qualities of climate, vegetation and land are the important factors affecting the prevalence of moulds in connection with a certain geographical location. Mould spores can disperse in the air with the wind or in combination of wind and rain [30]. The most frequent organisms implicated in mycotic mastitis were yeasts, especially of the genus *Candida. Candida spp.* are always present on the skin of the udder and teats, and could have entered the teat canal either due to inappropriate use of instruments or due to contaminated antibiotic preparations used for infusion [31].

With reference to Table 1, it is obvious that the most common isolated moulds from cows' nasal swabs were *Aspegillus sp.* (96%), *Fusarium sp.* (64%), *Mucor sp.*(8%), *Cladosporium sp.*(16%), *Alternaria sp.* (4%), *C.albicans* (24%), *C. tropicalis* (68%) and *Rhodotorula rubra*(12%). The present results coincide to some extent with that of Nilsson and Persson [32]; Maity and Deb [33]; Mahendra et al., [34]. Breeding factors such as animal housing, feeding on moldy hay and ventilation system or environmental factors such as temperature, wind and dew increase the odds of contracting the infection [35, 36].

Fungi can produce reproductive failure in animals either by direct infection of the genital system or by producing toxin metabolites (mycotoxins), which are subsequently ingested and absorbed. Moreover, mycotic abortion is the most important consequence of fungal infection of the genital tract [37].

From the obtained results in (Table 1), a total of 20 fungal species related to 8 genera were isolated from 25 samples of vaginal swabs collected from cases of cows suffered from abortion. The main recovered genera of fungi were *Aspegillus sp.* (80%), *Fusarium sp.* (16%), *Pencilium sp.*(32%), *Cladosporium sp.*(12%), *Alternaria sp.* (8%), *C.albicans* (40%), *C. tropicalis* (32%) and *Rhodotorula rubra*(3%). The results concur to some extent with the finding of Panangala *et al.*, [38]; Patgiri and Uppal [39]; Abdel-Rahman and Ibrahim [40] and [11] who isolated species of *Alternaria, Aspergillus,* and *Cladosporium, Candida, Fusarium* and *Pencillium* from vaginal swabs of repeat breeder cows. The predominance of dark-colored fungi may be explained by their higher resistance to dissication, Strong light and ionizing radiation than non-pigmented forms [41]. On the contradictory, *A. fumigatus* has been reported by Knudtson and Kirkbride [42] as the most common cause of mycotic placentitis. *Candida albicans* and *C. tropicalis,* were the most frequent yeast species recovered from cows' vaginal swab samples as yeasts have been implicated as causes of bovine reproductive problems, including abortion in cows and infertility in bulls [43, 44].

It is clear that the most common isolated mould from cows' feed were *Aspegillus* sp. (100%), *Fusarium* sp. (24%), *Mucor* sp.(28%), *Pencilium* sp.(52%), *Cladosporium* sp.(8%), *Alternaria* sp. (28%), *C.albicans* (80%), *C. tropicalis* (40%) and *Rhodotorula rubra*(32%). These findings were in agreement with the results of Arakawa [45] and Hassan et al. [4, 5, 6,11]. Also, it has been estimated that 25% of the world's crop production is contaminated with mycotoxins. Pusterla *et al.*, [46] suggested that the primary infection was attributed to inhalation of spores originating from moldy hay or soil.

In the present study, the current data in table (3) showed that, sera of twenty cases of diseased cattle outbreaks which suffered from loss of weight gain, low productivity, diarrhea, disturbance in fertility and sudden mortality of some cases, contained significant levels of aflatoxins and ZEAR. Meanwhile, 60% of cattle had the mean levels of aflatoxins (15.20±0.4 ppb) and zearalenon detected in 80% of cattle with the mean level of (62 ± 0.3) , respectively (Table, 2). Mycotoxins in sera cattle in Egypt in association with symptoms of toxicities were previously reported by Hassan [11]; Hassan et al. [2-6]. The effects of mycotoxins in human and animals were varied from carcinogenic; nephrotoxic and immunosuppressive health effects [47, 3-6]. Although the main route of human exposure to mycotoxins has been identified as the direct ingestion of contaminated cereals, grains and food of animal origin [47], there are many studies about whether the ingestion of meat, milk, and eggs originating from mycotoxin-exposed food-production animals is a significant pathway for mycotoxins among humans [48, 4, 5, 6,11]. In the feed samples were examined for fungal contamination and detection of mycotoxins. When, the used feed samples in breeding of these animals were subjected for detection of aflatoxins, the results revealed that the amounts of AFB1; OA; ZEAR.; T2; and FB1 were detected in (60%; 40%; 32%; and 44%) of feed samples, with the mean levels of $(55.0\pm1.50, 45.0\pm0.30)$ and 31.0 ± 0.2 and 40.0 ± 2.1 ppb), respectively (Table, 3). The significant levels of mycotoxins in the present feed samples and serum of diseased animals gave a large possibility that mycotoxins were responsible for the disease outbreak in animals. The Food and drug administration has established recommended maximum levels for aflatoxins in animal feed are 20 μg/kg of feed [49]. The permissible limits of aflatoxin for large ruminants were varied between 700-1000 ppm/ ten of feed which caused loss of weight gain, high food consumption and low feed efficiency [50]. Whereas, the O.A in each produced significant pathological changes in pregnant animal at the levels of 2.5 ppmof animal body weight for 5 days when given orally [51]. On the other hand the trichothecenes mycotoxins (T2) was produced such change at the doses as low as 10 ppm [52].

Therefore, the detected levels of mycotoxins were significantly over the permissible limits in feeds which may resulted of toxicosis due to continous feeding of toxiocated feed. The same findings were detected by many authors as [3-6], who detected such these diseased cases in association of significant high mycotoxin levels in feed and sera of diseased animals.

Therefore, the detoxification of toxic feed become a critical demand, ammonium hydroxide was the most safe chemical compound used for this purpose [1,3,4,5].

In this study, we demonstrated that the AFB1 and zearalenon in serum of cattle lead to a marked elevation in the serum enzymes activities of (AST, ALT and GGT) table (4) which indicated the hepatocellular damage, as previously reported [53,54] elevation could potentially be attributed to hepatic degeneration and subsequent leakage of enzymes into circulation after rupture of the plasma membrane and cellular damage [55, 56]. Such hepatic toxic effect of aflatoxin attributed to its active metabolite in liver as epoxide, Netke et al. [57] which covalently bind to DNA and may affect structural and enzymatic protein function [58]. Moreover the elevation of serum GGT activity suggested hepatic necrosis, thickness of bile duct and intrahepatic cholestasis [59]. The levels of serum urea and creatinine were significantly high in aflatoxicated groups as compared to healthy and that group of animals fed on the treated feed with ammonium hydroxid. This increase in concentrations in toxicated animals might be due to nephrotoxic action, which causes renal impairment by destruction of epithelial cells of proximal and distal convoluted tubules and alteration in tubular function [4, 60]. Also, if microbial protein synthesis in the rumen which is inhibited by mycotoxins, more free ammonia remains in the rumen, is absorbed into the blood, and is metabolized to urea, resulting in elevated blood urea concentrations and reduced hepatic fractional protein synthesis rates [61-63].

Mycotoxins in cattle and buffaloes inducing significantly decreased values in serum total protein, albumin, alpha globulin, beta globulin, and gamma globulin (table, 5). Decrease in serum globulin in mycotoxicated animals might be due to the adverse effect of mycotoxins on synthesis of total proteins and globulin. These results agree with Osuna and Edds [64] in pigs; Hassan and mogda [4] in quails and Madheswaran et al., [65] in chickens. Mycotoxins cause inhibition of DNA and protein synthesis as well as immunosuppressive due to the inflammation, cirrhosis of liver and kidneys [66,67]. At the same table (5) the globulin component showed drop in $\alpha 1$, $\alpha 2$, $\beta 1$ and $\gamma 1$ globulin in all the diseased animals, while increase γ 2globulin as compared with healthy animals. The results coincided with the tune of total proteins and albumin. This may be attributed to that AFB1 causes hepatotoxic, nephrosis, hemorrhages (liver and kidneys) [68]. In addition, mycotoxins has immunosuppressive effect inhibit nearly cellular and humeral immunologic reaction [68,2,5,4]. The decrease in serum level of globulin might be due to the adverse effect of mycotoxins on synthesis of total proteins and globulin [69, 70, 11].

In this study, we have shown that the mycotoxicated animals had significant increase in serum level of malondialdehyde (MDA) MDA. Mycotoxins particularly AFB1 are well known to generate free radicals, disturbing the antioxidant status and ultimately leading to oxidative stress and carcinogenesis [71]. Lipid peroxidation plays an important role in carcinogenesis [72] and may lead to the formation of several toxic products, such as malondialdehyde (MDE) and 4-hydroxynonenal. Also, AFB1 peroxide is a very reactive and unstable metabolite of AFB1 that will bind to cellular macromolecules like DNA, RNA, lipids and proteins, leading to lipid peroxidation and cellular injury. [73]. The hepatic damage is associated with an increase in the tissue malondialdehyde (MDA) level, an indirect index of lipid peroxidation [74].

The present study shows that serum NO level significantly increased in mycotoxicated cattle and buffaloes. It has been reported that elevated levels of lipid peroxidation stimulates host cells, mainly monocytes/ macrophages, to produce and release NO by induction of inducible nitric oxide synthase (iNOS) protein, resulting in cytotoxicity and DNA damage [75].

Our finding of low plasma retinol, and α -tocopherol, in toxicated cattle compared to apparently healthy and treated animals (table, 6) was shown in previous studies by Clarke *et al.*, [76], Vatassery et al., [77]. Aflatoxin causes oxidative stress by inducing the epoxide reactive ring, reducing the antioxidant defense systems of cells via depleting non enzymatic antioxidant system (vitamins and glutathione) and/or increasing susceptibility of cells to oxidative attack by altering the membrane integrity and fatty acid composition [78,79,80].

In conclusion, previously it were recorded that the fungal diseases affect upon the growth rate and health of human being and animals including aneamia , stunted growth, carcinogenic, tremorgenic, haemorrhagic, dermatitic, pulmonary edema, immunosuppressive and hormonal effects [4,5,,6,11]. This study, reported the dangerous effects of fungal diseases and their mycotoxins and its pollution of animal feed which resulted a significant losses in animal health and cause an important contributor to the country's economy in the form of meat, milk, wool and leather, with respect to the effects of environmental factors. Therefore, frequent testing program of the animal feeds and their environment for fungi and mycotoxin contamination were a critical demand to safe the animal and human health.

Fungal isolates	Milk of mastitic animal		Nasal swabs		Vaginal s	wabs	Feeds		
	No. of +ve	%	No. of +ve	%	No. of +ve	%	No. of +ve	%	
Aspergillus sp.	22	88	24	96	20	80	25	100	
A. candiduss	6	24	1	4	0	0	15	60	
A. clavatus	2	8	2	8	2	8	1	4	
A .flavus	20	80	20	80	13	52	18	72	
A. fumigatus	21	84	16	64	7	28	5	20	
A. niger	10	40	20	8	23	92	12	48	
A ochraceus	0	0	0	0	5	20	5	40	
Fusarium sp.	10	40	16	64	4	16	6	24	
F. oxysporum	10	40	1	4	0	0	6	24	
F. verticillioides	2	20	2	8	4	16	1	4	
Mucor species	0	0	4	8	0	0	7	28	
Penicillium sp.	2	8	0	0	8	32	13	52	
P. citrinum	1	4	3	12	0	0	0	0	
P. duclauxii	0	0	0	0	3	12	2	8	
P funiculosum	0	0	10	40	2	8	0	0	
P. islandicum	0	0	1	4	2	8	0	0	
P. oxalicum	1	4	0	0	4	16	11	44	
P.purpurogenum	0	0	0	0	0	0	2	8	
Cladosporium specie	2	8	4	16	3	12	2	8	
Al. alternata	1	4	1	4	2	8	7	28	
C. albicans	15	60	6	24	10	40	20	80	
C. tropicalis	6	24	17	68	8	32	10	40	
Rhodotorula rubra	4	16	3	12	3	121	8	32	

Table ([1]): Prevalence	of fungal	species in	clinical	samples of cattle
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- 25 samples of each were examined

Table (2): Determination of mycotxoins in serum of cattle.

Animal	Total No.	Aflatoxin B1			Ze	Learalenone				
		Positive cases	%	Mean levels ppb	Positive cases	%	Mean levels ppb			
Apparently healthy cattle	20	2	10	2.5±0.01	1	5	5.0 ± 0.0			
Diseased cattle	20	12	60	15.20±0.01	16	80	62.0±0.1			

Ppb : Part per billion

Table (3): Mycotoxins in feed consumed by cattle.

Mycotoxi	A	flato	xin B1	Ochratoxin A		Zearalenon			T-2			Fumonisin BI			
n in feeds	Posi	tive	Mean	Positive Mean		Positive		Mean	Positive		Mean	Posi	tive	Mean	
	No	%	of	No	%	of	No	%	of	No	%	of	No	%	of
			levels			levels			levels			levels			levels
			ppm			ppm			ppm			ppb			ppb
Animal	15	6	55±1.5	10	4	45±0.3	8	3	31±0.2	12	3	26±0.0	11	4	40±2.
feeds		0	0		0	0		2	0		6	2		4	1

- Fifty five samples of feed were examined.- Ppb : part per billion .

Chemical tests	Apparently	Diseased cattle	Treated cattle		
	healthy cattle	groups	groups		
AST	49.11±4.26	90.45 ± 7.21***	63.33±3.2*		
ALT	30.67 ± 3.84	51.34± 3.29***	43.33±3.8*		
ggt u/l	22.64 ± 1.15	64.18±4.49***	49.9±3.54*		
Urea	43.48±1.67	58.86±3.79**	48.65±1.6		
Creatinine	0.94±0.05	1.14±0.06*	0.96±0.96		

Table (4):Biochemical changes due to mycotoxicosis in cattle

*Results are expressed as means ± SEM (n =15), student 't' test

Table (5): Paterns of protein electrophoresis in diseased cattle.									
Paterns of protein	Apparently healthy cattle	Diseased cattle groups	Treated cattle groups						
T.protein	6.16 ± 0.09	5.42 ±0.16***	6.04±0.21						
Albumin	2.38 ±0.04	1.98 ±0.08***	2.26±0.07						
Alpha1	0.82 ±0.02	0.63 ±0.01***	0.72±0.03*						
Alpha2	0.35 ±0.03	0.29 ±0.01	0.34±0.05						
T-alph	1.17 ±0.04	0.92 ±0.01***	1.06±0.03*						
Beta1	0.66 ±0.02	0.56 ±0.01***	0.66±0.03						
Beta2	$0.40 \pm .02$	0.438 ± 0.02	0.48±0.06						
T.beta	1.06 ± 0.01	0.998 ±0.02*	1.14±0.05						
Gamma1	1.26 ±0.03	1.19 ±0.02**	1.29±0.13						
Gamma2	0.29 ±0.01	0.33 ±0.02	0.29±0.02						
T.Gamma	1.55 ±0.04	1.52 ±0.05	1.58±0.13						
T.globulin	3.78 ±0.05	3.43 ±0.09**	3.78±0.12						
A/G	0.62 ±0.01	0.57 ±0.01***	0.59±0.008*						

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*Results are expressed as means ± SEM (n =15), student 't' test

Table (6): Alteration of Levels of antioxidant in toxicated Buffaloes and cattle.

Antioxidants tests	Apparently healthy	Diseased cattle	Treated cattle		
	cattle	groups	groups		
Malondialdehyde	1.94±0.37	4.18±0.39***			
(MDA)(nmol/ml)			2.14±0.35**		
GSH(Glutathion)mmol/l)	12.4±6.8	18.2±5.7**	16.33±1.74		
Nitric oxide (N O)	35.17 ± 2.40	55.84 ± 6.50	48.01±4.21*		
(µmol/L)		**			
Vitamin A(ug/dl)	33.25± 2.44	28.67 ± 2.51	30.00±1.54		
Vitamin E(ug/ml)	14.54 ± 3.58	12.91±2.63	13.68±1.65		

*Results are expressed as means ± SEM (n =15), student 't' test

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