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Research Article

Study on Effect of Diatomaceous Earth (DAE) on Aflatoxin-Induced DNA Damage in Visceral and Lymphoid Organs in Broiler Chicken

Abstract

Background: Limited information exists concerning on the effect of diatomaceous earth (DAE) on aflatoxin-induced DNA damage in visceral and lymphoid organs.

Objectives: The present investigation is an attempt to detect the effect ability of Diatomaceous earth (DAE) in reducing the detrimental effects of aflatoxin (AF) in broiler diet was evaluated based on structural characteristic of DNA in liver, kidneys, heart, pancreas, thymus, spleen and bursa of Fabricius.

Materials and Methods: Three hundred and sixty healthy unsexed one day old broiler chicks were assigned to 9 groups comprising of control and treatment groups. DAE was supplemented @ 400 and 800 mg Kg⁻¹ of feed along with 0.5 and 1 ppm of AF Kg⁻¹ of feed for a period of 35 days. DNA fragmentation assay was conducted to detect changes in DNA.

Results: The DNA fragmentation in the toxin fed birds was severe in liver and lymphoid organs (thymus, spleen and bursa of Fabricius), followed by kidneys, heart and pancreas. The damage was more pronounced at 1 ppm in comparison to 0.5 ppm of dietary aflatoxin. The damage to DNA in most of the organs was reduced in birds of co-treatment groups fed with varying dosage of aflatoxin and DAE in the diet.

Conclusions: The present study showed that aflatoxin at graded doses induced marked DNA fragmentation indicating the genotoxic effect of aflatoxins. Addition of diatomaceous earth to aflatoxin mixed feed caused decreased DNA fragmentation in visceral and lymphoid organs.

Introduction

Aflatoxins (AF) are secondary metabolites and a class of mycotoxins produced predominantly by *Aspergillus flavus* and *A. parasiticus* [1]. The toxin occurs worldwide in feeds and feed stuffs resulting into severe economic loss to poultry and livestock industries in many countries around the world [2]. Studies have been related to negative effects of aflatoxin in broiler chickens including decrease in body weight gain, efficiency of feed utilization, liver damage, poor performance and immune responses. AF also caused pathologic alteration in important organs such as liver, kidneys and lymphoid organs [3]. The pathological change in broilers are characterized by hepatic lesions such as enlargement, paleness, fatty change, bile duct hyperplasia and periportal fibrosis [4]. Furthermore, the transmission of AF and its metabolites from feed to animal edible tissues and products, such as liver and eggs [5], becomes particularly important as a potential hazard for human health.

Producers and researchers desire to develop an effective detoxification technology dealing with the feed-borne toxin [6]. Approaches used included physical, chemical and biological treatment of contaminated feed and feed stuffs. A successful detoxification process must be economical and must be capable of eliminating all traces of toxin without leaving harmful residues and also should not impair the nutritional quality of the commodities [7]. In the last two decades several studies have been performed using adsorbent materials for detoxifying AF in contaminated feed and feed stuffs [8].

Several approaches to avoid contamination such as decontamination or remediation of feed and feedstuffs have been proposed [4]. A variety of adsorbents such as bentonite [9], zeolite [10], hydrated sodium calcium aluminosilicate [11], *Saccharomyces cerevisiae* [12], and activated charcoal [13], have been successfully used in detoxifying aflatoxins in contaminated feeds [14,15]. Adsorbent compounds utilized to ameliorate

aflatoxicosis in poultry diets includes aluminosilicates, bentonite, silicas, zeolite, quartz, etc have been evaluated for their ability to remove or diminish the adverse effect of mycotoxins in animal feed. These compounds must not be absorbed from the gastrointestinal tract and must have the ability to bind physically with chemical substances, precluding their adsorption [10]. The major advantages of adsorbents include cost effectiveness, safety and easy administration through animal feeds. However, no compound is completely effective in eliminating aflatoxin from feed.

Genomic DNA constitutes the total genetic information of an organism. The genomes of almost all organisms are DNA, the only exceptions being some viruses that have RNA genomes. Genomic DNA molecules are generally large and in most organisms are organized into DNA-protein complexes called chromosomes. The size, number of chromosomes and nature of genomic DNA varies between different organisms. Aflatoxins possess genotoxic potential which is mainly due to adduct formation with DNA, RNA and protein. This may be the most important product from the carcinogenic point of view.

Diatomite or Diatomaceous earth (DAE) is a kind of clay that consists of 90 per cent silicon dioxide. It is fine-grained, biogenic siliceous sediment and is available in large quantities at low cost [16]. DAE consists essentially of amorphous silica derived from opalescent frustules of diatoms resulting in an inert, light weight, highly porous, super-absorbent material, and has a fine porous structure with low density [17]. Modirsanei et al. [18], reported that diatomaceous earth increased the body weight gain, feed intake, and improved the feed conversion ratio as well as productive efficiency index. DAE also increased serum albumin levels in the birds that were subjected to AF supplementation.

Considering the beneficial properties of diatomaceous earth, the present study was undertaken to evaluate the efficacy of DAE in ameliorating aflatoxin induced patho-morphological changes in broilers.

Materials and Methods

Aflatoxins

Aflatoxin (AF) was produced on the rice using *Aspergillus parasiticus* (MTCC-2796) as per the method of Shotwell et al. [19], with some modifications and was quantified using thin layer chromatography.

Experimental birds and diet

Three hundred and sixty unsexed day-old healthy broiler chicks were procured from a reputed commercial hatchery and reared in battery cage system in experimental sheds with average temperature ranging from 27 to 31°C and relative humidity of 59% to 62% with 16:8±1 h Light : Dark cycle of intensity of 10 to 20 lux. All chicks were vaccinated on days 7 and 11 of age with the Lasota strain of Newcastle disease virus and Infectious bursal disease (intermediate strain) respectively.

Optimum conditions of brooding and management were provided to the birds throughout the period of experiment.

Toxin free and Diatomaceous earth (DAE) free Starter and finisher broiler feed was procured from Department of Poultry Science, Veterinary College, Hebbal, Bangalore-24, India as recommended by the National Research Council. Required quantity of cultured aflatoxin material was added to make the final concentration of aflatoxin in feed to be 0.5 ppm and 1ppm.

The birds were randomly divided into 9 groups, each comprising of 40 chicks. The different experimental group are as per the table 1.

All the birds were checked daily for the health and husbandry conditions. All the sanitary and hygienic precautions were strictly followed throughout the experiment. Prior permission of the Institute Animal Ethical Committee (IAEC) was obtained before the conduct of experiment. The birds were observed daily for clinical signs of aflatoxicosis characterised by dullness, poor growth, inappetence, diarrhoea and mortality (if any). A complete record of the daily mortality (if any) was also maintained.

Sample collection

At the end of the experimental period, the birds were euthanized by cervical dislocation and were subjected to detailed post-mortem examination. Representative tissue samples from liver, kidneys, pancreas, intestines, heart, thymus, bursa of Fabricius were collected, put on ice and immediately carried to the laboratory for DNA extraction.

Extraction of tissue DNA

The tissue DNA from liver, kidneys, heart, pancreas, thymus, spleen and bursa of Fabricius of birds belonging to all the groups was extracted as per the standard protocol, using the DNeasy® Blood and Tissue kit procured from Qiagen, Inc., (USA).

Protocol for DNA extraction

- a) The tissues (≤ 10 mg spleen or ≤ 25 mg other tissue) were cut into small pieces, and placed in 1.5 ml microcentrifuge tube. 180 μ l Animal Tissue lysis (ATL) buffer and 20 μ l proteinase K were added and mixed by vortexing for 4-5 seconds, and incubated at 56°C until completely lysed. Vortexing was done occasionally during incubation.

Table 1: Experimental design for various treatment groups.

Groups	Aflatoxin (ppm)	DAE(ppm)	No. of Birds
I	0	0	40
II	0	400	40
III	0	800	40
IV	1	0	40
V	0.5	0	40
VI	1	400	40
VII	1	800	40
VIII	0.5	400	40
IX	0.5	800	40

- b) 200 µl of Lysis buffer (AL) was added and mixed thoroughly by vortexing by pulse-vortexing for 5–10 seconds.
- c) Two hundred microlitre of absolute ethanol (99%) was added to the sample and mixed by pulse-vortexing for 15 seconds.
- d) After mixing, the microcentrifuge tube was briefly centrifuged at 3000 rpm for 1 minute to remove the drops from inside the lid.
- e) The mixture from the above step (including the precipitate) was carefully applied to the QIAamp mini spin column along with 2 ml collection tube without wetting the rim. The cap was closed and centrifuged at $\geq 6000 \times g$ for 1 min. Spin column was placed in a new clean 2 ml collection tube and collection tube containing the filtrate was discarded.
- f) 500 microlitre of wash buffer (AW1) was carefully added without wetting the rim of the spin column.
- g) The cap was closed and centrifuged at $\geq 6000 \times g$ for one minute and the collection tube was discarded.
- h) The spin columns were carefully opened and 500 µl of wash buffer (AW2) was added without wetting the rim of the spin column.
- i) The cap was closed and centrifuged at $20,000 \times g$ for 3 minute.
- j) QIAamp mini spin column was placed in a clean 1.5 ml microcentrifuge tube and 200 µl elution buffer (AE) or distilled water was carefully added and incubated at room temperature for one minute. Finally, the DNA was eluted at $\geq 6000 \times g$ for 2 minute.

Determination of purity and yield of the DNA samples

The purity and concentration of the extracted tissue DNA was estimated by UV spectrophotometry. An aliquot of 20 µl of DNA sample was dissolved in 0.98 ml of sterile distilled water. The diluted DNA was transferred into 1ml microcuvette and the optical density (OD) was checked at 260 nm and 280 nm in a UV spectrophotometer. Sterile distilled water was used as blank. The ratio of OD at 260/280 nm was calculated. A ratio of 1.7 to 1.9 was considered pure. The concentration of the DNA was estimated by the equation: $1 \text{ OD } 260 \text{ nm} = 50 \text{ µg/ml}$ of DNA. Further, the purity of the DNA sample was checked by electrophoresis on 1.8 per cent agarose gel.

Protocol for DNA confirmation and fragmentation study By Agarose Gel Electrophoresis

For electrophoresis, 1.2 % agarose was used and it was carried out as follows.

- a) The edges of a clean, dry, gel casting tray sealed at both the ends using adhesive tape. An appropriate comb was placed to form a sample slot in the gel.

- b) Agarose solution was prepared by dissolving required quantity of agarose in proportionate volume of 1X Tris-Borate-Ethylenediaminetetraacetic acid (TBE) buffer and melted in a microwave oven for 1 min.
- c) Once the molten gel cooled, 0.5 µg of ethidium bromide was added and mixed thoroughly by gentle swirling.
- d) Warm agarose solution was then poured into the gel casting tray avoiding formation of air bubbles and allowed to solidify.
- e) Once agar gel solidified, a small amount of Tris-acetate-Ethylenediaminetetraacetic acid (TAE) buffer was poured on the top of the gel to remove the comb. Then the buffer was poured off and the tape was removed.
- f) The gel casting tray was mounted in the electrophoresis tank and the TAE buffer was added just enough to cover the gel to a depth of one mm.
- g) 5 µl of DNA was mixed with 1/6th volume of 6X gel loading dye and slowly loaded into the slots of submerged gel using a micropipette.
- h) The gel tank was closed with the lid and electrical leads were attached so that the DNA would migrate towards the anode.
- i) The electrophoresis was carried out at 5V / cm at room temperature (RT) until the bromophenol blue dye migrated appropriate distance through the gel.
- j) Following electrophoresis, DNA bands were visualized at 300 nm wavelengths using a UV transilluminator and the images were captured by using Gel Doc XR (Bio Rad, USA) for further interpretation.

Results

The results of quantification of isolated DNA from various organs are expressed in ng/µl by Nanodrop method using spectrophotometer has been summarized in tables 2,3. The fragmentation of DNA in various tissues of broiler chickens in different treatment groups at the end of experiment (35th day) has been analysed and the same is presented as below.

The DNA fragmentation in AF toxin fed groups was observed in all the organs examined such as liver (Figure 1, Table 4), kidneys (Figure 2, Table 5), thymus (Figure 3, Table 6), bursa (Figure 3, Table 6) and spleen (Figure 4, Table 7).

AF feeding caused severe damage to DNA of liver and lymphoid organs namely thymus, spleen and bursa of Fabricius; followed by kidneys and the least damage was observed to the DNA of heart and pancreas. The damage was more pronounced at 1 ppm (Group IV) in comparison to 0.5 ppm (Group V) of dietary aflatoxin.

The DNA from various tissues of birds fed diet free of aflatoxin (Group I) and DAE alone to toxin free feed (Group II and III) showed no damage to the DNA.

Table 2: DNA quantification in various visceral organs of different groups by Spectrophotometer (Nanodrop method).

GROUPS	LIVER		KIDNEY		HEART		PANCREAS	
	Nucleic Acid Conc. (ng/μl)	260/280	Nucleic Acid Conc. (ng/μl)	260/280	Nucleic Acid Conc. (ng/μl)	260/280	Nucleic Acid Conc. (ng/μl)	260/280
BLANK	-0.6	0.66	-0.6	0.66	-0.6	0.66	-0.6	0.66
I	202.6	1.65	352	1.77	172.4	1.92	888.4	1.83
II	464.9	1.64	445.5	1.79	168.8	1.89	632.7	1.71
III	126.4	1.74	451.6	1.79	100.8	1.98	1369.4	1.8
IV	518.6	1.78	114.9	1.67	245.9	1.77	2976.2	1.87
V	298.6	1.82	251.2	1.79	179	1.89	585.9	1.68
VI	610.4	1.79	723.9	1.82	790.7	1.97	1086.9	1.82
VII	346	1.93	1212.9	1.89	332	1.85	969.8	1.84
VIII	300.1	1.9	239.2	1.79	348.4	1.95	378.3	1.8
IX	204.3	1.79	328.9	1.71	2589.2	1.92	755.4	1.69

Table 3: DNA quantification in various lymphoid organs of different groups by Spectrophotometer (Nanodrop method).

GROUPS	THYMUS		SPLEEN		BURSA OF FABRICIUS	
	Nucleic Acid Conc. (ng/μl)	260/280	Nucleic Acid Conc. (ng/μl)	260/280	Nucleic Acid Conc. (ng/μl)	260/280
BLANK	-0.6	0.66	-0.6	0.66	-0.6	0.66
I	3751.3	1.89	2168.8	1.91	703.1	1.76
II	590.7	1.78	1052.4	1.92	2536.1	1.87
III	5909.5	1.88	223.3	1.87	6223.6	1.9
IV	7809.7	1.81	1528.6	1.9	2485.1	1.79
V	3794.3	1.87	495.2	1.89	789.8	1.85
VI	1558.9	1.73	892.9	1.85	509.1	1.86
VII	3881.1	1.84	1822.9	1.89	471.3	1.92
VIII	4454.3	1.9	2904.7	1.88	1130.8	1.89
IX	12377.8	1.86	274.2	1.85	749.9	1.89

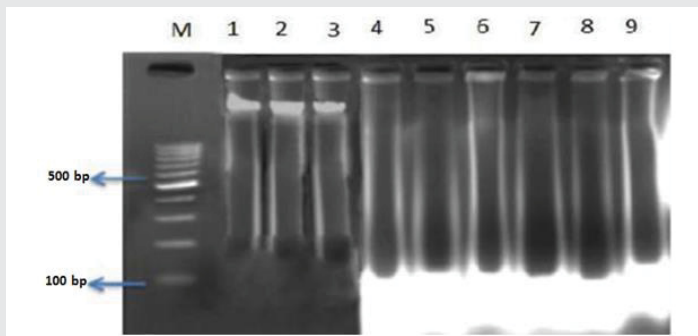


Figure 1: DNA fragmentation in liver at the end of fifth week in different treatment groups.

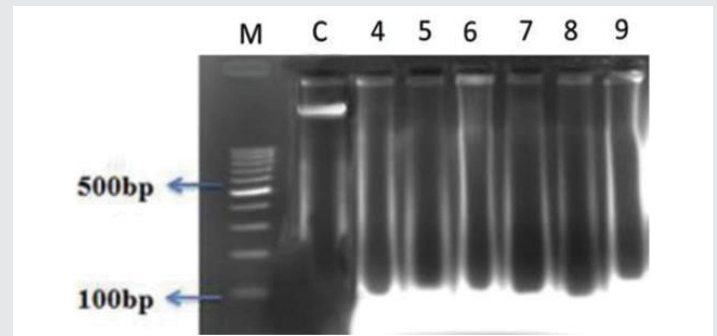


Figure 2: DNA fragmentation in kidneys at the end of fifth week in different treatment groups.

Table 4: Degree of DNA fragmentation in liver at the end of fifth week in different treatment groups.

Lanes	Particulars	Level of Fragmentation in base pair (bp)	Degree of DNA Fragmentation
M	100 bp Ladder		
1	Control group	DNASTreak/Smear	No fragmentation
2	Control+400 mg DAE	DNASTreak/Smear	No fragmentation
3	Control+800mg DAE	DNA Streak/Smear	No fragmentation
4	Aflatoxin 1ppm	<100	2+
5	Aflatoxin 0.5 ppm	<100	2+
6	Aflatoxin 1ppm+400mg DAE	<100, >1200	2+
7	Aflatoxin 1 ppm+800mg DAE	<100, >1200	1+
8	Aflatoxin 0.5 ppm+400mg DAE	<100, >1200	2+
9	Aflatoxin 0.5ppm+800mg DAE	>100, >1200	1+

Table 5: Degree of DNA fragmentation in kidneys at the end of fifth week in different treatment groups.

Lanes	Particulars	Level of Fragmentation in base pair (bp)	Degree of DNA Fragmentation
M	100 bp Ladder	Kidney	Kidney
1	Control group	DNA Streak/Smear	No fragmentation
2	Control+400 mg DAE	DNASTreak/Smear	No fragmentation
3	Control+800mg DAE	DNASTreak/Smear	No fragmentation
4	Aflatoxin 1ppm	DNASTreak/Smear	2+
5	Aflatoxin 0.5 ppm	<100	1+
6	Aflatoxin 1ppm+400mg DAE	DNA Streak/Smear	No fragmentation
7	Aflatoxin 1 ppm+800mg DAE	<100	1+
8	Aflatoxin 0.5 ppm+400mg DAE	>100	1+
9	Aflatoxin 0.5ppm+800mg DAE	100-200	1+

The damage to DNA in most of the visceral and lymphoid organs was significantly reduced in bird of co-treatment groups fed 0.5 and 1ppm of aflatoxin and DAE @ 400 and 800 mg/kg feed (Group VI to IX). The birds fed with DAE @ 800 mg/kg feed and aflatoxin (0.5 and 1 ppm) showed reduced damage to DNA in comparison to broilers fed with DAE @ 400 mg/kg feed and aflatoxin @ 0.5 and 1 ppm.

Discussion

Genomic DNA constitutes the total genetic information of an organism. The genomic DNA in cells can be subjected to injury following interaction due to exposure to toxicants. Mycotoxins, especially the aflatoxins are known to cause interaction with the cellular total genomic DNA [20,21]. This interaction can result in small to large changes in genomic DNA evident as DNA fragmentation [22], similar to the findings of the present study.

The fragmentation assay of DNA extracted from heart, liver, kidneys, pancreas, thymus, spleen and bursa of birds fed with different dietary levels of aflatoxin in the present study revealed higher damage at 1 ppm in comparison to 0.5 ppm. Following aflatoxin treatment, severe damage caused to DNA of liver and lymphoid organs, followed by kidneys and the least damage was observed in DNA of heart and pancreas in comparison to negative control groups. The liver is a target organ for AF toxicity as it is the site where aflatoxins undergo bioactivation to reactive 8,9-epoxide, which then binds to DNA and proteins [23], causing wide spread damage to the hepatic tissue. In addition, aflatoxins have direct tropism for any of the primary or secondary lymphoid organs resulting into either the organ necrosis or atrophy or depletion of specific subpopulations of the lymphoid cellular components [24].

Similar to the findings in the present study, Shebl et al. [25], reported the higher mean percentages of DNA fragmentation

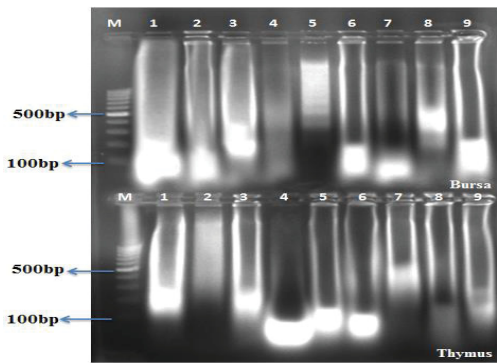


Figure 3: DNA fragmentation in bursa of Fabricius and thymus at the end of fifth week in different treatment groups.

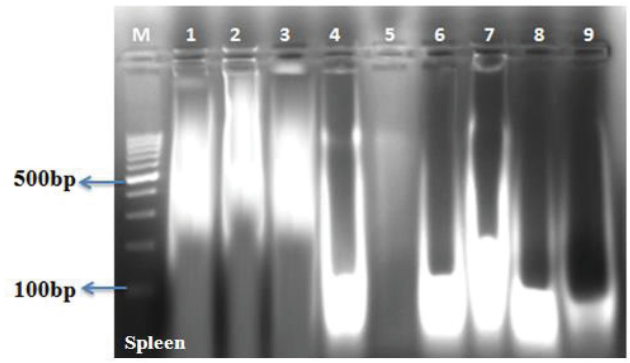


Figure 4: DNA fragmentation in spleen at the end of fifth week in different treatment groups.

Table 6: Degree of DNA fragmentation in bursa of Fabricius and thymus at the end of fifth week in different treatment groups.

Lanes	Particulars	Level of Fragmentation in base pair (bp)		Degree of DNA Fragmentation	
		Bursa	Thymus	Bursa	Thymus
M	100 bp Ladder				
1	Control group	DNA Streak/Smear	DNASTreak/Smear and 150-300	No fragmentation	No fragmentation
2	Control+400 mg DAE	DNA Streak/Smear	DNA Streak/Smear	No fragmentation	No fragmentation
3	Control+800mg DAE	DNASTreak/Smear and 200-400	DNASTreak/Smear and 150-300	1+	No fragmentation
4	Aflatoxin 1 ppm	DNASTreak/Smear and >1200	<100	1+	1+
5	Aflatoxin 0.5 ppm	400,500,600,700	>100	4+	1+
6	Aflatoxin 1 ppm+400mg DAE	100-200	<100	1+	1+
7	Aflatoxin 1 ppm+800mg DAE	<100	200-500	1+	1+
8	Aflatoxin 0.5 ppm+400mg DAE	200-500	200	1+	1+
9	Aflatoxin 0.5ppm+800mg DAE	100-250	200-300	1+	1+

Table 7: Degree of DNA fragmentation in spleen at the end of fifth week in different treatment groups.

Lanes	Spleen -Particulars	Level of Fragmentation in base pair (bp)	Degree of DNA Fragmentation
M	100 bp Ladder		
1	Control group	DNA Streak/Smear	No fragmentation
2	Control+400 mg DAE	DNA Streak/Smear	No fragmentation
3	Control+800mg DAE	DNA Streak/Smear	No fragmentation
4	Aflatoxin 1 ppm	>100 and 1000	2+
5	Aflatoxin 0.5 ppm	DNA Streak/Smear and 1000	1+
6	Aflatoxin 1 ppm+400mg DAE	>100	1+
7	Aflatoxin 1 ppm+800mg DAE	>200	1+
8	Aflatoxin 0.5 ppm+400mg DAE	<100	1+
9	Aflatoxin 0.5ppm+800mg DAE	<100	1+

and increase in the frequency of micronucleated cells in liver cells following feeding of aflatoxin @ 211.88 µg / kg feed to the broiler in comparison to negative control groups and opined that AFB₁ was selective inhibitor of DNA synthesis in mammalian cells and can induce DNA adducts, induce mutations by intercalating to DNA by forming adduct with guanine moiety in the DNA. Similar findings were also reported by earlier workers following mycotoxicosis [26,27].

Al-Terehi [28], observed higher degree of DNA fragmentation in liver and spleen in comparison to kidney and blood in female albino rats following feeding of aflatoxin for a period of 2 weeks. The DNA damage observed in various organs could be attributed to AFB₁- DNA adduct formation at cellular level, as suggested by Tolliver and Robbins [29]. In addition, oxidative stress to rats following exposure to AFB₁ can also induce DNA lyses by forming (8-oxo d G) in liver cells [30].

Aflatoxin possess genotoxic potential through the mixed-function oxidase system to a number of hydroxylated metabolites and to aflatoxin 8, 9 epoxide which binds to DNA, forming covalent adducts [31], and disturbs DNA replication causing genetic alteration [32]. The adduct formation occurs preferably with guanine resulting in AFB₁-N-7 guanine adduct responsible for mutagenesis in AFB₁ treated cells [33].

DNA fragmentation test in the present study clearly indicated that aflatoxin could not damage the nucleus of cells in different organs of DAE supplemented birds (Group VI to IX), which might be explained by the strong adsorption of AF to diatomaceous earth and thereby preventing the entry of AF into circulation [27]. Similar to the findings of present study, Shebl et al. (2010) [25], also reported that the addition of clay material (DAE) either alone or in combination with aflatoxin reduced the mean percentages of DNA fragmentation in different organs.

The effectiveness of compound in sequestering one mycotoxin does not mean an equal ability to sequester other mycotoxins due to chemical complexity of these toxins. Each of the mycotoxin has different functional groups; thus, the binding capacity of an adsorbent depends on its chemical and physical properties and its relation with the physical structure of the target mycotoxins. Thus, the physicochemical differences among the adsorbents used in the studies mentioned above could explain the higher or lower efficacy among them. However, the ability of the toxin binder to bind mycotoxins depends on other factors such as pH, molecular arrangement and its geographic region of origin [34]. Natour and Yousef [35], reported significantly higher *in-vitro* adsorption ability of DAE to aflatoxin, which is directly proportional to the number of diatom valves. Diatomaceous earth being a powerful natural adsorbent can effectively adsorb the mycotoxins through its polar ends [36]. In addition, DAE has a small mass (0.5–0.8 g/cm³), high porosity and high content of silicon responsible for the high adsorption capacity [37]. *In-vitro* study showed that DAE has high (94.71 %) ability to absorb AF from the feed at pH 6.5 [38]. The normal pH of the chicken intestinal tract contents is 5.7–6.0 in the duodenum/jejunum, 6.3–6.4 in the ileum/rectum and finally up to pH 7.0 or higher in the caecum (Denbow, 2000) [39]. Considering the correlation between the pH and ability of mycotoxin binder in *in-vitro* studies, higher

binding ability of DAE to the aflatoxins can plausibly be expected at the pH of 6–7 in the intestinal tract of chicken to reduce the absorption and systemic availability of this mycotoxin.

In conclusion, the incorporation of DAE in the diet during the period of exposure to AF in the present study could considerably reduce the toxic effects of aflatoxin. This study highlights the protective effects of DAE, which might be due to its capability of specific chemisorption of aflatoxin in gastrointestinal tract, which reduces AF bioavailability by formation of aflatoxin-DAE complex followed by excretion through droppings/faeces [40,41].

The compounds like clay, zeolite minerals and DAE are structurally and functionally diverse; they vary considerably from source to source and may not have equal affinities and capacities for binding of aflatoxin and other mycotoxins. Thus these adsorbents should be rigorously tested one by one and thoroughly in *in-vivo* conditions, paying particular attention to their effectiveness and safety in sensitive animal/avian models and their potential for harmful interactions. Similarly, generalisations should be avoided for all potential mycotoxins detoxifying agents, as adsorbing compounds can differ in efficacy even within the same category. Considering the results of present study and earlier work done on effect of different levels of DAE on aflatoxin further studies employing the broader perspectives seems to be necessary to determine whether lower levels of DAE in broilers diet will be effective in controlling or preventing the occurrence of aflatoxicosis in chicken.

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