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## Fungal Contaminant of Poultry Feed in Basrah, Iraq

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**Abstract:** A total of 180 samples of concentrated poultry feed were collected from different broilers, broiler breeders, layers farms and local market of poultry in Basrah province. Feed samples were collected during the period from Sep. 2014 to Apr. 2015. About 10 - 30 representative samples of 1 kg were collected from several locations. They were cultured on Potato dextrose agar (PDA) and malt extract agar (MEA) and then sub cultured on Sabouraud dextrose agar (SDA). Seven genera were recovered from 180 samples of poultry feed. The most genera which recovered were *Aspergillus* spp. frequency (Fr) 62.77% - Relative density (RD) 52.03%, followed by *Penicillium* spp. (Fr 47.77% - RD 17.01%) and they were the predominant genera isolated from poultry feed. *Fusarium* spp. isolates were less frequency and relative density (Fr.1.66%, RD 2.11%). The most frequently isolated *Aspergillus* spp. was *Aspergillus flavus* (Fr 65.48%) and had the most RD (27.55%), followed by *Aspergillus niger* (Fr. 58.40%, RD14.23%), the less occurrence of *Aspergillus* spp. was *Aspergillus parviticus* (Fr.1.76%, RD0.89%).

### 1. Introduction

A large quantity of animal feeds purchased from abroad and with the increased emphasis on animal resources, this amount is expected to increase substantially in the coming years. It is necessary to have quality control of the feed because the feed can have a great effect on the weight and mortality of birds which prepared as starter and finisher diets for poultry [1; 2]. Diets for poultry consist largely from grain, nutritional supplements such as concentrated protein, soybean meal, vitamin and mineral [3; 4].

The storage conditions are necessary for safe feed, so weather extremes, unsuitable storage practices, and improper of feed condition can cause feed contamination, especially with mycotoxins [5]. Poultry feed is frequently contaminated by mycotoxins [6;7]. Poultry feed it is more susceptible to fungal growth during processing, therefore identification of fungi with the ability to produce mycotoxins is essential [8]. There are many clinical signs as a result of mycotoxins effected with low concentration were always observed in different broiler, broiler breeder, it is appeared reduced weight gain, anorexia, feed conversion efficiency, egg production and egg shell quality [9]. Molds can grow and produce mycotoxins in preharvest or during storage, transport, processing or feeding and during these periods, humidity and temperature play an important role in the fungal growth and mycotoxins contamination [10]. In wet feeds, higher moisture levels allow mold growth if oxygen is available [11]. Feeds containing more than 12-15% moisture suitable to grow fungi and Because most molds are aerobic, high moisture concentrations can eliminate air and prevent mold growth [12]. The conditions are most suitable for mold growth and for mycotoxin production are not necessarily the same [13]. More than 100.000 fungal species are considered as natural contaminants of agricultural and food products.



The present study was aim to investigate the occurrence ,distribution and frequency of fungal contaminant of poultry feed.

## 2. Materials and methods

### 2.1. Collection of samples

A total of 180 samples of poultry feed pellet were collected from different breeders broiler farms and local markets in Basrah province. Feed samples were collected during Sep. 2014 to Apr. 2015, and stored for 2-3 days in sterile containers at room temperature (22-25°C).Then, they were prepared for fungal isolation and identification [2].

### 2.2. Isolation and identification of fungi

Twenty gram poultry feed samples were suspended with 180 ml of saline solution (0.85% Sodium Chloride) in addition to 0.05% Tween 80 (polyoxyethylene sorbitan monooleate) on a horizontal shaker for 30 min. to liberate the spores from fruiting bodies and to break the spore clumps [4]. 0.1 ml of suspension was inoculated on PDA and MEA media [15; 16]. The distinct colonies were stained on a slide using lactophenol cotton blue and lacto-fuchsin, then morphological characteristics of fungal isolates were described by microscope [17; 18]. The colony color and conidia morphology were investigated. Each colony type was counted for individual cfu/g counts and was recorded [19].

### 2.3. Statistics

The frequency (Fr.) and relative density (RD) of isolation genus and species were calculated as follows [20; 21 ; 22]:

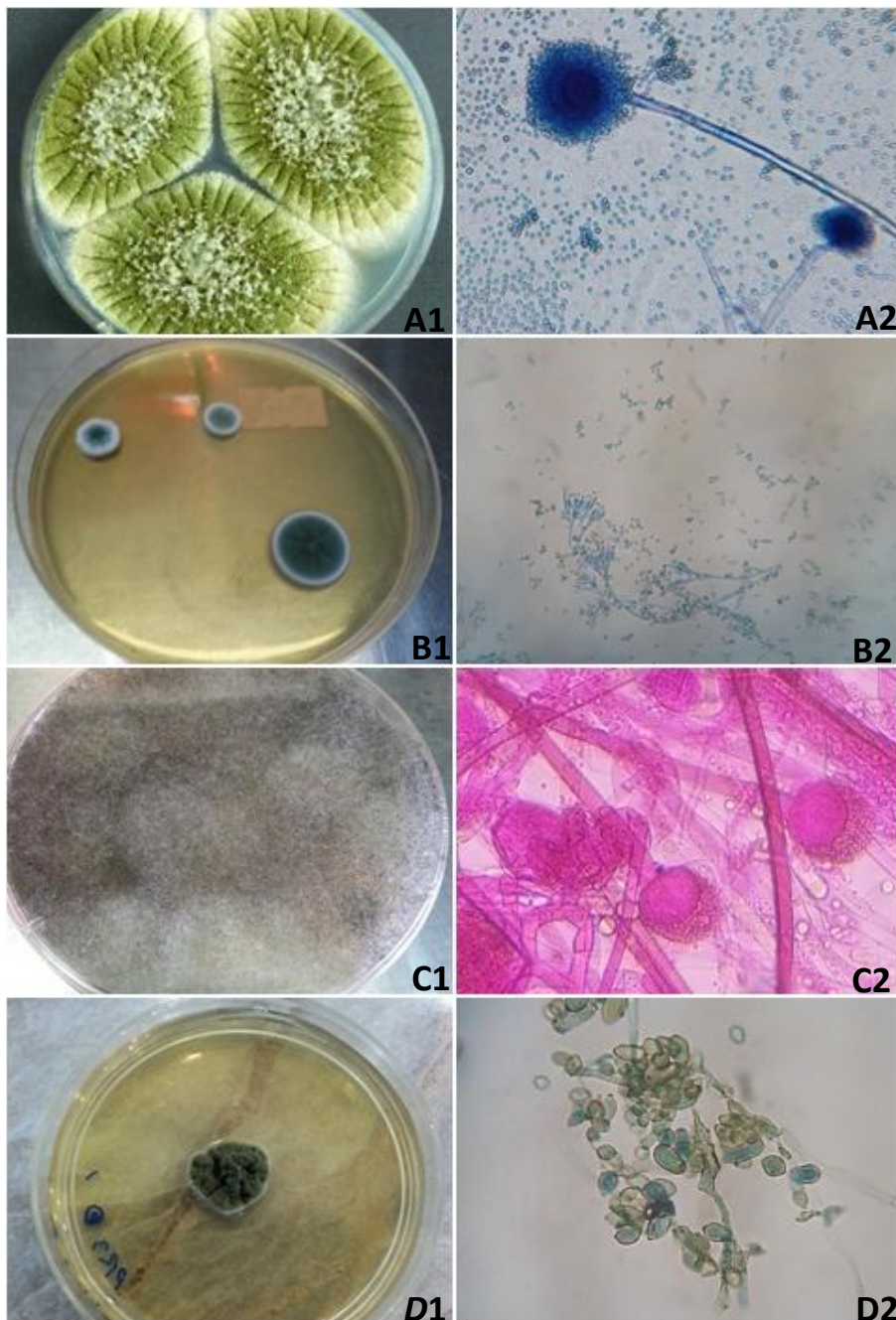
$$\text{Fr. (\%)} = \frac{\text{samples number with a genus or species}}{\text{samples total number}} \times 100$$

$$\text{RD(\%)} = \frac{\text{isolates number of a genus or species}}{\text{Total number of fungal isolates}} \times 100$$

## 3. Results

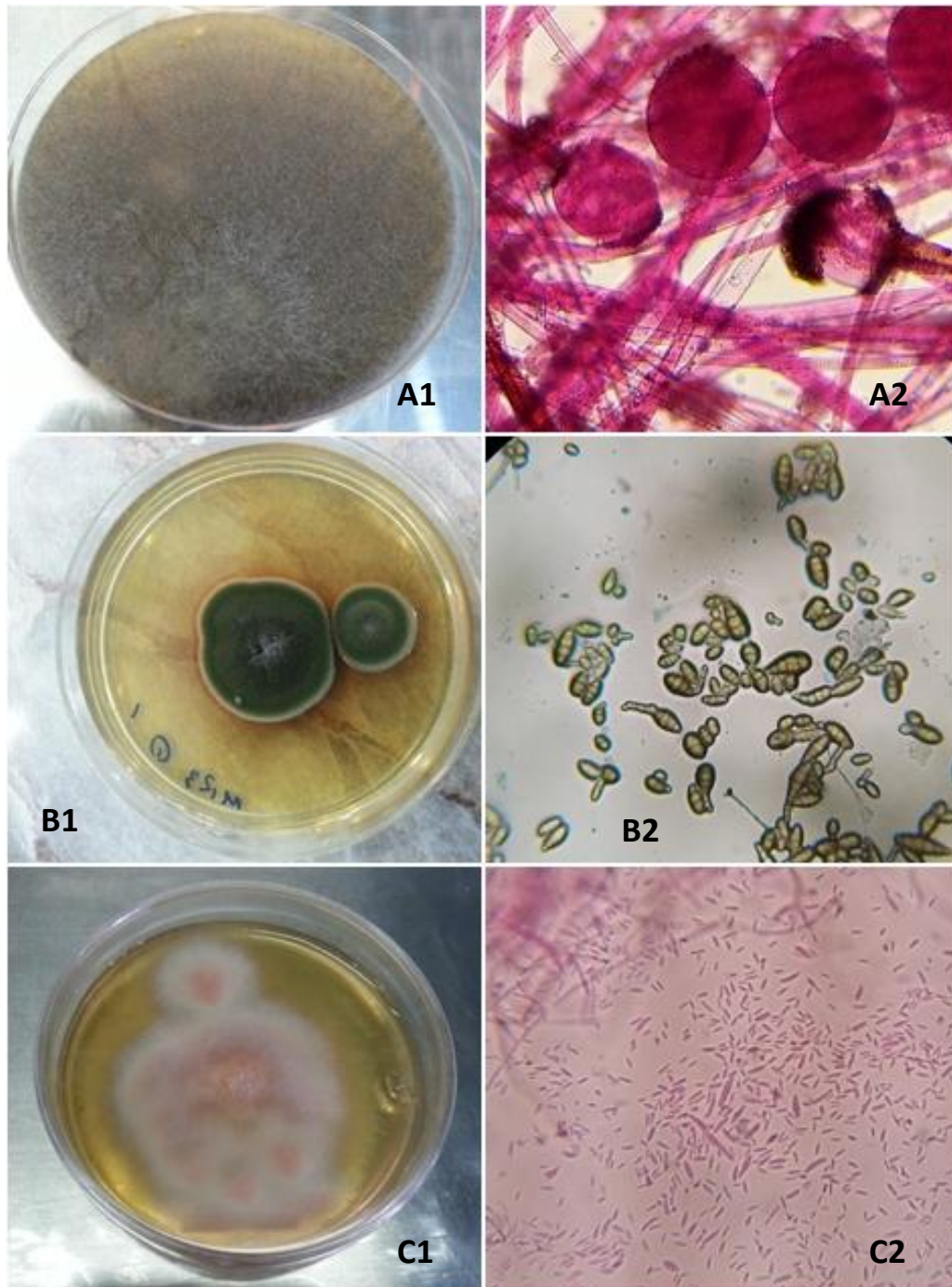
### 3.1. Fungal isolation

In the present study, about seven fungal genera and forty species were identified from 180 samples of poultry feed. Several species were identified in each sample. The morphological characteristics of fungal isolates were described under the microscope. The colony color and conidial morphology were investigated. The most important recovered genera of fungi were *Aspergillus* spp., *Penicillium* spp., *Rhizopus* spp., *Cladosporium* spp., *Mucor* spp., *Alternaria* spp. ,and *Fusarium* spp. (figure 1 and 2). There were 9 *Aspergillus* spp. Recovered; *A. flavus*, *A. niger*, *A. fumigatus*, *A. terreus* , *A. flavipes*, *A. carbonarius*, *A. ochraceus*, *A. candidus* and *A. parasiticus*.



**Figure 1: The isolated molds genera from poultry feed on PDA medium. A: *Aspergillus* sp., B: *Penicillium* sp., C: *Rhizopus* sp., D: *Cladosporium* sp. (1): In culture, (2): Microscopically, 40X.**





**Figure 2: The isolated molds genera from poultry feed on PDA medium. A: *Mucor* sp., B: *Alternaria* sp., C: *Fusarium* sp.. (1): In culture ,(2): Microscopically , 40X .**  
The total fungal counts cfu/g were ranged from  $5 \times 10^1$  -  $2.1 \times 10^6$  of feed sample, with an average of  $1.5 \times 10^5$  cfu/g of a sample ( tables 1) .

**Table 1: Range and average count cfu/g of recovered molds genera from poultry diet samples.**

Genus	Total mold count (CFU/g)	
	Range	Mean values
<i>Aspergillus</i> spp.	2.2X10 <sup>5</sup> - 2.1X10 <sup>6</sup>	1.1X10 <sup>6</sup>
<i>Penicillium</i> spp.	7.1X10 <sup>3</sup> - 8.0X10 <sup>4</sup>	4.3X10 <sup>4</sup>
<i>Rhizopus</i> spp.	4.2X10 <sup>3</sup> - 2.1X10 <sup>4</sup>	1. 2X10 <sup>4</sup>
<i>Cladosporium</i> spp.	2.4X10 <sup>3</sup> -2.5X10 <sup>3</sup>	2.4X10 <sup>3</sup>
<i>Mucor</i> spp.	1.5X10 <sup>3</sup> -2.0X10 <sup>3</sup>	1.7X10 <sup>3</sup>
<i>Alternaria</i> spp.	6.0X10 <sup>1</sup> - 6.3X10 <sup>1</sup>	6.1X10 <sup>1</sup>
<i>Fusarium</i> spp.	5.0X10 <sup>1</sup> - 5.6X10 <sup>1</sup>	5.3X10 <sup>1</sup>

The frequency (Fr.) and relative density (RD) recorded the highest value which belonged to *Aspergillus* spp., while *Fusarium* sp. was identified as a low value of Fr. and RD (Table 2). The most frequent mycotoxigenic fungi from 180 samples were those from the genus *Aspergillus* spp.. This genus recovered from 113 samples (Fr. 62.77%) also with most RD= 52.03% with a range of 2.2X10<sup>5</sup> - 2.1X10<sup>6</sup> cfu/g, and a mean value of 1.7X10<sup>6</sup> cfu/g, followed by *Penicillium* spp. which recovered from 90 samples (Fr.47.77%), also the RD17.01% with a range of 7.1X10<sup>3</sup> - 8.0X10<sup>4</sup> cfu/g, and a mean value 4.1X10<sup>4</sup> cfu/g, *Rhizopus* was recovered from 86 samples with Fr. 50% and RD10.01% , had a range 4.2 X 10<sup>3</sup> - 2.1X10<sup>4</sup> cfu/g, with mean value 1.6X10<sup>4</sup> cfu/g. *Fusarium* spp., was found as a low ratio, it was recovered from 7 samples (Fr. 1.66%) RD 2.11% with a range of 5X10<sup>1</sup>- 5.6X10<sup>1</sup> cfu/g and of a mean value of 7.8X10<sup>1</sup> cfu/g.

**Table 2: Frequency and relative density of recovered mold genera from poultry feed samples.**

Genus	Fr. of Positive samples	Fr. %	RD%
<i>Aspergillus</i> spp.	113	62.77	52.03
<i>Penicillium</i> spp.	90	47.77	17.01
<i>Rhizopus</i> spp.	86	50	10.01
<i>Cladosporium</i> spp.	21	11.66	6.23
<i>Mucor</i> spp.	21	4.44	2.55
<i>Alternaria</i> spp.	8	3.88	2.24
<i>Fusarium</i> spp.	7	1.66	2.11

The total *Aspergillus* spp. counts cfu/g were ranged from  $0.3 \times 10^2$  -  $1.5 \times 10^5$  of feed samples, with an average  $7.5 \times 10^4$  cfu/g sample (table 3).

**Table 3: Range and average count cfu/g of recovered *Aspergillus* spp. from poultry feed samples**

<i>Aspergillus</i> spp	Total mold count CFU/g	
	Range	Mean values
<i>A. flavus</i>	$1.1 \times 10^4$ - $1.5 \times 10^5$	$8.0 \times 10^4$
<i>A. niger</i>	$5.8 \times 10^3$ - $7.4 \times 10^3$	$6.6 \times 10^3$
<i>A. fumigatus</i>	$1.7 \times 10^3$ - $3.0 \times 10^3$	$2.3 \times 10^3$
<i>A. terreus</i>	$1.2 \times 10^3$ - $2.4 \times 10^3$	$1.8 \times 10^3$
<i>A. flavipes</i>	$0.5 \times 10^2$ - $1.0 \times 10^2$	$7.5 \times 10^2$
<i>A. carbonarius</i>	$0.4 \times 10^2$ - $0.7 \times 10^2$	$5.5 \times 10^1$
<i>A. ochraceus</i>	$0.4 \times 10^2$ - $0.7 \times 10^2$	$5.5 \times 10^1$
<i>A. candidus</i>	$0.3 \times 10^2$ - $0.4 \times 10^2$	$3.5 \times 10^1$
<i>A. parasiticus</i>	$0.2 \times 10^2$ - $0.4 \times 10^2$	$0.3 \times 10^1$

The frequency and RD recorded the highest value which belonged to *A. flavus*, while *A. parasiticus* was identified as a low value of Fr. and RD (Table 4). The most predominant *Aspergillus* species recovered from 113 samples were *A. flavus*, recovered from 74 (Fr. 65.48 %) with RD 27.55 % and with range,  $1.1 \times 10^4$  -  $1.5 \times 10^5$  and had a mean value of  $8.6 \times 10^4$  cfu/g, followed by *A. niger*, recovered from 66 samples (Fr. 58.40%) and RD 14.23%, with a range  $5.8 \times 10^3$  -  $7.4 \times 10^3$  and a mean value  $9.5 \times 10^3$  cfu/g, *A. fumigatus*, recovered from 19 samples (Fr. 16.81%) with RD (7.60%), recorded with range  $1.7 \times 10^3$  -  $3.0 \times 10^3$  cfu/g, and mean value  $3.2 \times 10^3$  cfu/g, *A. parasiticus*, were recovered as low percentage and cfu/g, it was recovered from 2 samples with Fr. 1.76 % and with RD 0.89%, which recorded a range  $0.2 \times 10^2$  -  $0.4 \times 10^2$  cfu/g, with mean value  $0.4 \times 10^1$  cfu/g.

**Table 4: Frequency and relative density of recovered *Aspergillus* spp. from poultry feed samples**

<i>Aspergillus</i> spp.	Fr. of Positive samples	Fr. %	RD%
<i>A. flavus</i>	74	65.48	27.55
<i>A. niger</i>	66	58.40	14.23
<i>A. fumigatus</i>	19	16.81	7.60
<i>A. terreus</i>	12	10.61	0.50
<i>A. flavipes</i>	9	7.96	2.13
<i>A. carbonarius</i>	8	7.07	2.40
<i>A. ochraceus</i>	6	5.30	2.12
<i>A. candidus</i>	3	2.65	1.34
<i>A. parasiticus</i>	2	1.76	0.89

#### 4. Discussion

Total fungal counts in this study were ranged from  $5 \times 10^1$ -  $2.1 \times 10^6$  of feed sample, with an average  $1.5 \times 10^5$  cfu/g sample and were considered as a high contaminant to poultry feed as compared with GMP [23] which demonstrated that the fungal propagules were useful indicators to determine a quality of feeds hygienic, which should not over a value of  $1 \times 10^4$  cfu/g. These results agree with those reported by Bragulat et al., [24] from Spain, in Argentina [25], in Brazil [8], and similar to the study in Iraq [2].

This study reported different results of that in Argentina [26], and in Brazil [27], that which found the level of cfu/g in their study about 103 cfu/g. They deal with samples in different ways from this study in collection place, and the quantity of samples used in homogenizing because they collected monthly during one year from factories directly from the production line (after processing) and used ten grams of each sample for homogenizing also they incubated under cold white and black fluorescent lamps in a 12/12 h photoperiod for 7 days. The processing way could cause a decrease of different microorganisms during storage from New Zealand [28], and most of the fungi are slow-growing under these conditions and observations may be made after 20 days [29] in Ethiopia. In this study, the samples were collected from farms and the local market of poultry, then 20 g of samples were used in homogenizing then incubated in the dark incubator, therefore the mean of cfu/g of this study was higher 105.

The genera of *Aspergillus* was the predominant isolated from poultry feed in this study with a mean value of  $(1.7 \times 10^6)$ cfu/g (Fr 62.77% - RD 52.03%), because temperature during collection was a range of 37-47°C and considers the optimal for growth of *Aspergillus* [31], followed by *Penicillium* with a mean value of  $4.1 \times 10^4$  cfu/g (Fr 47.77% - RD 17.01%) and *Rhizopus* had mean value of  $1.6 \times 10^4$  cfu/g (Fr.50% -RD10.1%). Similar results which were found in Spain [31; 24], in Argentina [25, 26], in Brazil [32], and in Iraq [2], they reported that the *Aspergillus* genera were the most frequent. While it differs from the study in Serbia [27] who reported the most frequent genus isolated was *Penicillium* followed by *Aspergillus* and *Fusarium*. While two reports in Serbia [10, 26] they confirmed that the *Fusarium* is the predominant genera followed by *Penicillium* and *Aspergillus*, because the location of Serbia is in the moderate continental climate belt, geographical factors are of superior importance for the occurrence of *Fusarium* and the most frequently isolated fungi contaminating feedstuffs, vegetables, cereals, and fruits are from the *Fusarium* [33; 34], also the *Fusarium* was reported as the most genus recovered from poultry feed, followed by *Eurotium*, *Penicillium*, and *Aspergillus*, they collected their samples from the region in Argentina suitable to permanent of *Fusarium* growth [26]. The *Aspergillus* spp. in this study was confirmed there were nine species recovered and the *A.flavus* was the most frequently and RD isolated *Aspergillus* with the mean value of  $8.6 \times 10^4$  cfu/g and (Fr. 65.48%, RD 27.55%), followed by *A.niger*, was reported with mean value of  $9.5 \times 10^3$  cfu/g and with Fr. 58.40%, RD 14.23%, and *A.fumigatus* with the mean value of  $3.2 \times 10^3$  cfu/g and with Fr. 16.81%, RD 7.60%. The highest dominance of *A.flavus* in this study agrees with previous studies [26, 27, 35, 36], but different to that in Pakistan who found that the most frequently isolated *Aspergillus* were *A. niger* followed by *A. flavus*, as a result of high humidity and high temperature might be responsible for a higher frequency of *A. niger* in poultry feeds compared with other species of *Aspergillus* [22]. Such climate was found in the area of this study which has been found very rich in different species of fungi [37; 38; 39; 40; 41].

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