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Research Article

Mycological Evaluation and Mycotoxin Contamination of Swine and Poultry Feed-Shelf Life Assessment in Makurdi, Nigeria



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Abstract

Mycotoxin production as a result of mycotic contamination of animal feeds presents a major concern to public health. It therefore becomes imperative to generate data on the mycological contamination and potential mycotoxins production with the aim of determining suitable shelf life for swine and poultry feed. For a period of eight weeks, a total of 25 swine feed samples including five samples each of Dusa (Grain husk), Soya beans, Bone meal, Brewer's residue (Burukutu) and Rice offal and 20 poultry feeds samples including five samples each of Broiler starter, Broiler finisher, Chick mash and Layer mash were collected on a weekly basis from the University of Agriculture Makurdi (UAM) animal farm and tested immediately for Moisture Content (MC), Total Mold Count (TMC), Fungal Isolation Frequency (FIF) and Relative Density (RD). The relative abundance of mycotic load from swine feed include Aspergillus species (42.6%), Fusarium species (36.4%), Alternaria and Acremonium species (29.6% each) in bone meal, brewers residue and rice offal respectively while for the poultry feed Aspergillus (38.1%) and Penicillium species (33.3%) in starter and finisher, Penicillium, Rhizopus and Curvularia had 27.6% in the chick mash with 20.0% of Curvulariain the layers mash. Mycotoxins detected included aflatoxin B1 and B2, ochratoxins A and B, patulin, sterigmatocystin, zeralenone and citrinin. Our results show the presence of mycotoxins in feed vis-a-vis deterioration in feed quality thus reducing shelf-life.

Keywords: Aspergillus; Mycotoxin; Penicillium; Poultry; Sterigmatocystin; Swine

Abbreviations: MC: Moisture Content; TMC: Total Mold Count; FIF: Fungal Isolation Frequency; RD: Relative Density; MAE: Malt Extract Agar; TLC: Thin Layer Chromatography; LOD: Limits of Determination; LOQ: Limits of Quantification

Introduction

Mycotoxin production as a result of mycotic contamination of animal feeds presents a major concern to public health due to the possible transmission of these toxic metabolites to meat, milk and animal products [1]. Mycotoxins include Aflatoxins (AFLs), OchratoxinA (OTA), Sterigmatocystin (EST), Fumonisins (FBS) and Zearalenone (ZEA) which are produce by Aspergillus spp., *Penicillium spp.* and *Fusarium spp.* in the tropics [2]. The production of these toxigenic secondary metabolites frequently occurs during the processing and storage of feed when environmental conditions such as moisture content and temperature becomes favourable for the proliferation of spoilage fungi thereby contaminating the entire food-chain from agricultural cultures to the plates of consumers [3,4]. Ingestion of mycotoxins causes unconcealed mycotoxicosis which is often associated with impairment of the immune system leading to a decrease in intake of feed, animal agility and reproductive ability. Mycotoxins have also been

incriminating in oncogenesis, mutagenesis and teratogenesis [5,6]. Owing to the indigenous cultural practices which encourage a high consumption of swine produce and a relatively high humidity in the study area there is a major risk factor for mycotoxin intoxication. It is imperative therefore, to generate data on the mycological contamination and potential mycotoxins production with the aim of determining suitable shelf life for swine and poultry feed in the study area.

Materials and Methods

Source of samples

A total of 25 swine feed samples including Dusa (Grain husk) (5), Soya beans (5), Bone meal (5), Brewer's residue (Burukutu)(5) and Rice offal (5) and 20 poultry feeds samples including Broiler starter (5), Broiler finisher (5), Chick mash (5) and Layer mash (5) were collected on a weekly basis for a period of eight weeks

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from the University of Agriculture (UAM) animal farm in Benue State, Nigeria. These primary samples were homogenized to form a composite sample and afterwards quartered to get 15.0g laboratory samples and tested immediately for Moisture Content (MC), Total Mold Count (TMC), Fungal Isolation Frequency (FIF) and Relative Density (RD). They were preserved at 4oC for mycotoxin analysis.

Moisture content analysis

Moisture content in grams was determined using the dry weight determination method. One (1g) gram of feed sample was weighed using a sensitive weighing balance and afterwards put into a dry air oven at 1000C to remove available moisture for sixty minutes. Results obtained in grams from the respective samples were recorded.

Mycobiota determination

Enumeration of mycotic propagules was done using pour plating technique [7]. Ten grams of each sample was emulsified and aseptically transferred into ninety 90ml of 0.1% peptone water. The mixture was shaken vigorously to dislodge possible fungal propagule in the sample. Serial dilutions to 10-3 were carried out for each sample using sterile pipette. Aliquots of one 1ml of 10-3 dilution were inoculated unto Potatoe Dextrose Agar supplemented with 0.05mg/ml chloramphenicol to inhibit opportunistic bacteria in triplicates. Fungal plates were kept at room temperature for seven days. Plates with 10-100 colonies were recorded as CFU/g. Distinctive Colonies were afterwards transferred to Malt Extract Agar (MAE) and incubated at 30 °C for microscopic and macroscopic identification. Taxonomic identification of the different genera and species were carried according to Pitt and Hocking [7] and Klich [8]. The isolation frequency of each genus and relative density of each species were calculated as follows [1]:

$$F = (ns/N) \times 100$$
; RD (%)=(ni/NI)×100

Where F is the Isolation frequency, ns is the number of samples in which a particular genus occurred, N is the total number of samples, RD is the relative density, ni is the number of isolates of a particular genus and NI is the total number of fungal isolates obtained.

Mycotoxin production

Three genera which included Aspergillus, Fusarium and Penicillium were tested for mycotoxin production according to methodology described by Soares and Rodrigues-Amaya [9] with slight modifications. Conidial suspension for the test organisms were prepared and afterward (100µl) of each strain was inoculated into 250ml conical flasks containing 30ml of YES culture medium (2% Yeast Extract, 15% Sucrose) which were grown in stationary phase and incubated at 30 °C for 30 days in the dark. Aliquots (1ml) of each culture was then mixed with 1ml chloroform and centrifuged at 4000g × for 10min. The chloroform phase was transferred to a clean vial, evaporated to dryness and re-suspended in 0.5ml methanol and rapidly subjected to thin layer chromatography (TLC). The analyses were carried out in a saturated vat with the following solvent system: methanol, ethyl acetate,

formic acid (60:40:0.5). Mycotoxins were detected under UV light and quantified by comparison against toxin standards (λ: 256 and 365nm). The Limits of Determination (LOD) and Quantification (LOQ) were 1 and 2 μ g L⁻¹.

Statistical analysis

Data were entered into Microsoft excel spread sheet and later transferred into SPSS 21 (SPSS Inc., Chicago, IL, USA) statistical software for analysis. To ensure that data were normally distributed, the transform menu function was used transform data log10(x+1). Analysis of Variance was then used to pre determine significant difference between means at P< 0.05.

Results

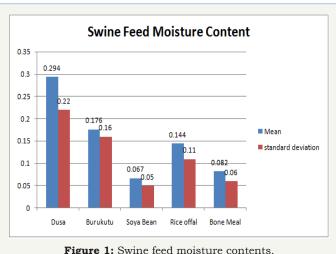


Figure 1 Highlights the moisture content in the swine feed samples analyzed. Grain husk 'dusa' had the highest moisture content of (0.294±0.22) followed by brewers residue 'burukutu' (0.176 ± 0.16) , rice offal (0.144 ± 0.11) , bone meal (0.082 ± 0.06) and soya beans (0.067±0.05) respectively. Poultry feed samples indicated highest moisture content of (0.214±0.13) in chick mash, (0.164±0.11) in layer mash, (0.131±0.12) in the broiler starter and (0.086±0.07) in the broiler finisher feed samples respectively (Figure 2).

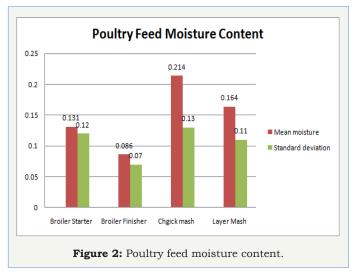


Table 1 Highlights the relative abundance of the mycotic load within the swine feed samples analyzed. Aspergillus species had a relative abundance of 42.1% and 36.4% in bone meal and brewers' residue 'burukutu' feed samples respectively, followed by Fusarium species with an abundance of 36.4% in the brewers

residue feed sample while Alternaria and Acremonium species had an abundance of 29.6% each in the rice offal feed however, an abundance of 8.3% was observed of Fusarium and Alternaria species in the grain residue 'dusa' feed sample.

Table 1: Mycotic Load in Swine Feed Samples (x10-3 CFU/g).

Fungal Isolates	Dusa		Burukutu		Soya bean		Rice offal		Bone meal	
	RD (%)	Average	RD (%)	Average	RD (%)	Average	RD (%)	Average	RD (%)	Average
Aspergillussp	8(22.2)	8.38±1.68	8(36.4)	8±2.44	7(25.9)	4.86±1.34	0(0)		8(42.1)	8.5±1.77
Rhizopussp	8(22.2)	2.75±0.46	0(0)	0	3(14.3)	2.57±0.57	3(11.1)	2.67±0.57	0(0)	0
Mucorsp	8(22.2)	4.25±2.37	0(0)	0	3(14.3)	2±0.0	8(29.6)	6.63±2.2	0(0)	0
Peniciliumsp	6(16.6)	5.17±1.94	6(27.3)	6.83±0.75	7(25.8)	5.14±2.34	0(0)		3(15.8)	1.67±1.115
Fusariumsp	3(8.3)	4.67±3.05	8(36.4)	3.5±1.41	0(0)	0	0(0)		0(0)	0
Aternariasp	3(8.3)	4.33±1.53	0(0)	0	7(25.9)	5.57±1.27	8(29.6)	10±2.20	8(42.1)	8.13±2.0
Acermoniumsp	0(0)	0	0(0)	0	0(0)	0	8(29.6)	7.0±2.4	0(0)	0
Total	36 (100)	5.03±2.62	22(100)	6.05±2.62	27(100)	4.56±1.90	27 (100)	7.30±3.02	19(100)	7.26±3.03

Table 2: Mycotic Loadin Poultry Feed Samples (x10-3 CFU/g).

Fungal Isolates	Broiler starter		Broiler Finisher		Layer starter		Layers mash	
	RD(%)	Average	RD (%)	Average	RD (%)	Average	RD (%)	Average
Aspergillussp	8(38.1)	8.25±1.83	8(33.3)	8.88±2.03	0(0)	0	0(0)	0
Penicilliumsp	8(38.1)	6.63±1.30	8(33.3)	6.38±1.85	8(27.6)	9.63±2.55	8(26.7)	8.37±1.68
Rhizopussp	5(23.8)	3.20±0.45	8(33.3)	3.13±0.84	8(27.6)	3.13±0.83	8(26.7)	3.37±1.06
Curvulariasp	0(0)	0	0(0)	0	8(27.6)	3.63±0.52	6(20.0)	5.5±1.51
Fusariumsp	0(0)	0	0(0)	0	5(17.2)	7.4±1.95	0(0)	0
Mucorsp	0(0)	0	0(0)	0	0(0)	0	8(26.6)	2.75±0.46
Total	21(100)	6.43±2.39	24 (100)	6.13±2.87	29 (100)	5.52±3.45	30 (100)	4.97±2.59

Table 2 & 3 Highlights the relative abundance of the mycotic load within the poultry feed samples as follows *Aspergillus* and *Penicillium* species recorded a relative abundance of 38.1% and 33.3% in the broiler starter and broiler finisher feed samples

respectively, Penicillium and Rhizopus and each had a relative abundance of 27.6% in the chick mash while, abundance of 23.8% of Rhizopus in the broiler starter and 20.0% of Curvularia in the layers mash feed samples were observed.

Table 3: Anova.

		Sum of Squares	df	Mean Square	F	Sig
SB	Between Groups	78.968	2	39.484	19.646	0
	With in Groups	36.175	18	2.01		
	Total	115.143	20			
	Between Groups	133	2	66.5	24.234	0
FB	Within Groups	57.625	21	2.744		
	Total	190.625	23			
	Between Groups	265.416	3	88.472	32.61	0
SL	Within Groups	67.825	25	2.713		
	Total	333.241	28			
ML	Between Groups	154.217	3	51.406	32.799	0
	Within Groups	40.75	26	1.567		
	Total	194.967	29			

Figure 3 Highlights isolation frequencies of the mycotic genera isolated from the swine feed samples. *Alternaria* species had an isolation *frequency* of 100% in the bone meal, rice offal and soya beans feed samples. Fusarium species was predominantly

isolated from the 'dusa' and brewer's residue 'burukutu' feed samples. Aspergillus species was predominant in the grain husk 'dusa', brewers residue 'burukutu 'and soya beans feed samples. Penicillium species predominated the soya beans and brewers residue 'burukutu' samples while Rhizopus and Mucor species predominanted the grain husk 'dusa' feed samples.

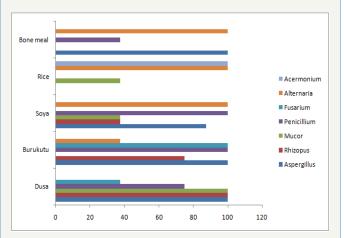


Figure 3: Isolation frequencies of different genera isolated from respective swine fee.

Figure 4 Highlights isolation frequencies of the mycotic genera isolated from the poultry feed samples Mucor, Rhizopus and Penicillium species where predominantly isolated from the layer mash feed samples with and isolation frequency of 100%. However, a predominance of Curvularia was indicated in the chick mash. Broiler feed samples indicated infestation by Rhizopus, Penicillium and Aspergillus species. Figure 5 Highlights a section of mycotoxins isolated based on their wavelengths. Mycotoxins detected included aflatoxin B1 and B2, ochratoxins A and B, patulin, sterigmatocystin, zeralenone and citrinin.

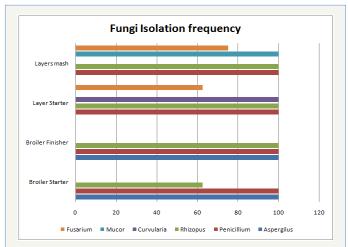


Figure 4: Isolation frequencies of different genera isolated from respective poultry feed.

Discussion

The quality of animal feed largely affects output. Productivity in animal breeding is often related to nutritional content of the feed therefore, emphasis is placed on improving nutrient quality of feed by addition of a variety of nutrient rich compounds and combination or rotation of different kinds of feed thus, it is not uncommon to have a form of feed rotation practice when breeding animal such as swine in the study environment. However, fed quality in terms

of feed health and safety is a vital aspect to ensure a productive output Gremmel, 2004. Investigations in this study indicated that the swine and poultry feed samples were infested with a varied mycotic genera which included Aspergillus spp., Penicillium spp., Fusarium spp., Rhizopus spp., Alternaria spp., Acermonium spp., Mucor spp. and Curvularia spp. concurring with reports by Zain et al. [4]. Swine and poultry feed samples recorded highest mycotic abundance of 42.6% and 38.1% respectively of Aspergillus species. Magnoli et al. [10], reported relatively high isolation frequencies for Aspergillus flavus, Lebars-Baiely et al. [11] suggested that the genus Aspergillus possess thermophilic and thermo resistant characters thus, their tendencies to proliferate under storage conditions in tropical climates. Investigations from the study also showed that alflatoxins B1 and B2, ochratoxins A and B and Sterigamatocystin were detected at respective wavelengths (Figure 5).

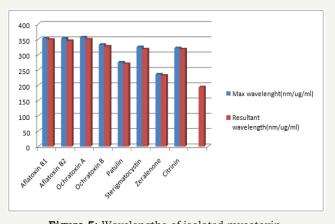


Figure 5: Wavelengths of isolated mycotoxin.

These mycotoxins have been implicated in a number of health conditions such as carcinogenesis, hypersensitivity reactions. Aspergillus spp. are also known to produce spores so their presence in animal feed is a concern to feed handlers as their spores have been implicated in respiratory inflammations, lung cancers and a reduced immunity; Rosa [2] reported ochratoxins produced by Aspergillus niger in substrates. Penicillium species was observed to proliferate in the poultry feed samples with an abundance of 33.1%, the abundance of this species would account for the detection of mycotoxins such as citrinin and patulin. These mycotoxins have been implicated in Mycotic abundance of 26.6% as observed of Mucor spp. in layer mash samples and 42.1% was observed for Acremonium in the bone meal samples, Rippon [12] reported that Mucor and Clasdosporium species may cause mycotic abortions and allergic reactions in livestock and humans owing to respiratory and systemic transmissions. Fusarium spp. with an abundance of 36.1% was observed in the swine feed sample; the fungus has been associated with invasive mycosis and are associated with opportunistic infections in both man and animals Khosravi & Gupta et al. [13] reported a dominance of 6% of Fusarium in feed samples, Gbore & Egbunike [14] reported FB1 levels of 5000µgkg ¹ reduces sperm production. Fusarium has been implicated in the production of the mycotoxins zeralenone which have been reported to influence oestrogenic activity in swine, sheep and cattle [15]. In

conclusion, results indicated relative mycotic load in both swine and poultry feed samples.

In spite of this, statistical analysis showed no significant difference was observed in the mycotic abundance within an eight week period. Arguments to extend sampling period may not hold sway because of high turnover in consumption of feed with respect to low feed production capacity as opposed to developed communities where there may be longer period of storage due to availability of feed raw material and mechanized production and surplus output. However, the heterogeneous form of feeding creates room for introduction of contamination as evident in the presence of mycotoxins in feed thus affecting shelf-life, it is safe to infer that there is deterioration in quality of feed within the sampling period in term of feed health and safety which is vital to productive output as it is necessary to ensure physiological functions and defense against diseases. To this end, it becomes imperative therefore to ensure that grains for feed are harvested at maturity, damage to produce is avoided at all stages of feed preparation, maintenance of even temperature during storage and use of extra caution when dealing with high moisture grain.

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