

INTEGRATIVE TOXICOLOGICAL EVALUATION OF AFLATOXIN B₁ EXPOSURE
IN FISCHER 344 RATS AND BIOMARKER BASED EXPOSURE ASSESSMENT OF
AFLATOXIN B₁ IN HUMAN POPULATIONS

by

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ABSTRACT

Aflatoxin B₁ (AFB₁) is a potent toxic and carcinogenic mycotoxin, which can adversely affect the immune system in animals and humans. Traditional toxicological evaluation of AFB₁ has focused on alternations in the histopathological, serum biochemical, and immunohistological aspects, an integration of molecular biomarkers (such as AFB₁-lysine adduct, AFB-Lys) data into these changes might provide useful information for a risk assessment purpose. Biomarkers based AFB₁ exposure or risk assessment is currently favored compared to food survey based exposure assessment. A rapid non-antibody based method for measuring serum AFB-Lys adduct in animals and humans was developed. The solid-phase-extraction based high performance liquid chromatography (HPLC) method has a limit of detection of 0.4 pg/mg albumin and recovery rates of 72.0-94.4%. The elimination kinetics of serum AFB-Lys and the toxicities of AFB₁ in F344 rats were further studied using single- (50-1000 µg/kg b.w.)

and repeated-dose (5-75 $\mu\text{g}/\text{kg}$ b.w./day) designs. Several kinetic parameters of serum AFB-Lys were determined: the peak time (4 h), the half-life (2.31 days) and a conversion ratio of 1.12-1.98% at 24 h, as estimated by a physiologically based pharmacokinetic model. A linear increase in the adduct level was found only after repeated low doses (5-25 $\mu\text{g}/\text{kg}$). Liver damage, bile duct proliferation and necrosis, were most prominent at 3 days after a single dose (≥ 25 $\mu\text{g}/\text{kg}$), correlating with clinical biochemistry changes (ALT and AST) and liver glutathione S transferase placental form positive (GST-P⁺) hepatocytes formation. Repeated treatment induced concurrent appearance of liver GST-P⁺ foci and bile duct proliferation after 3 weeks. One-week treatment decreased the percentages of splenic CD4⁺ and CD8⁺ T cells as well as their production of IL-4 and IFN- γ (≥ 25 $\mu\text{g}/\text{kg}$), and TNF- α production by CD3⁻CD8a⁺NK cells (75 $\mu\text{g}/\text{kg}$). Five-week treatment increased IFN- γ production by CD4⁺ cells (25 $\mu\text{g}/\text{kg}$), and TNF- α production by NK cells (75 $\mu\text{g}/\text{kg}$) and inhibited IL-4 production by CD4⁺ and CD8⁺ cells, suggesting an inflammatory response. These results provide an integrative toxicopathological evaluation of AFB₁ exposure in F344 rats. Human exposures were also assessed based on the measurement of serum AFB-Lys levels. The rank from high exposure to low is: West Africa countries (Ghana and Burkina Faso) > Southeast Countries (China and Malaysia) > Uganda > Haiti > United States. Currently, AFB₁ exposure remains prevalent in West Africa and Southeast Asia (detection rates > 90%), which warrants urgent and effective interventions. These data may facilitate human aflatoxin exposure assessment and improve our understanding of how AFB₁ can affect immune response.

INDEX WORDS: Aflatoxin, aflatoxin B₁ lysine, biomarker, toxicity, liver, spleen, exposure

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CHAPTER 1

INTRODUCTION

Problem Statement

Aflatoxins represent a group of naturally occurring mycotoxins that have long been recognized as hazardous contaminants of food, especially in peanuts and corn. Among the naturally occurring aflatoxins (B₁, B₂, G₁, G₂), aflatoxin B₁ (AFB₁) is the most potent hepatotoxic and genotoxic, and has been categorized as a known human carcinogen (Group I) by the International Agency for Research on Cancer (IARC) ¹. Acute exposure to high levels of AFB₁ via the diet causes disease (aflatoxicosis) and death in animals and humans ². Chronic exposure to low levels of aflatoxins is one of the major risk factors in the etiology of human hepatocellular carcinoma (HCC) in several regions of Africa and Southeast Asia ³. In addition, AFB₁ has also been shown to be an anti-nutritional agent in animals and humans and also a potential immunotoxic agent in humans ^{4, 5}, which may aggravate infectious diseases in the developing world, such as HIV infection ⁶. Aflatoxin contamination in food remains a serious burden in the developing countries where poverty and contaminated food supplies present a major and persistent challenge. Human exposure can occur by consumption of aflatoxin contaminated commodities and the products derived from them as well as from tissues, eggs, and milk (AFM₁) of animals that have consumed contaminated feeds ⁷. Occupational exposure, as noted in poultry workers through inhalation, has gained much

concern^{8, 9} and deserves caution. Therefore, accurate assessment of global aflatoxin exposure is urgently needed.

Biomarkers are cellular, biochemical, or molecular alterations that can be accurately measured in biological media, such as human tissues, cells or fluids, to indicate the exposure, effect or susceptibility in individuals. The knowledge of the toxicology and metabolism of aflatoxins leads to the validation of several biomarkers that have gained wide application in human exposure assessment and efficacy evaluation of mitigation strategies for reducing aflatoxin exposure. These biomarkers include urinary AFB₁-N⁷-guanine adduct, aflatoxin M₁, AFB₁-mercapturic acid, and serum AFB₁-albumin adduct. AFB₁-albumin adduct, or more specifically, AFB₁-lysine adduct, has a relative longer biological half life than other urinary biomarkers and serves as a good marker for chronic exposure for up to several months. Chemical-specific biomarkers can be used to identify stages in the development of the toxic effects/disease of environmental agents and to provide important information for critical regulatory, clinical and public health problems.

Currently available methods for measuring serum AFB₁-lysine adduct consists of enzyme linked immunosorbent assay (ELISA), immunoaffinity column based HPLC method, radioimmunoassay (RIA), and the isotope dilution mass spectrometry (IDMS) method. The results of antibody-based assays are influenced by the specificity and the sensitivity of the antibodies used. Since the major AFB-albumin adduct was identified as the AFB₁-lysine adduct, the measurement of the latter has been widely used in human studies. A non-antibody based method that doesn't require sophisticated mass spectrometry¹⁰ seems promising towards initiation of a global network for monitoring

aflatoxins exposure or assessing efficacy of intervention strategies to reduce aflatoxin exposure.

Traditional toxicological evaluation of AFB₁ has focused on alternations in the histopathological, serum biochemical, and immunohistological aspects¹, an integration of molecular biomarkers (such as AFB₁-lysine adduct) data into these changes might provide useful information for a risk assessment purpose. Such data are relatively scarce, especially the correlation between changes in biomarker and toxicity in a time course.

The overall objective of my research is to develop a non-antibody method for measuring serum AFB₁-lysine adduct, to take an integrative toxicological evaluation of AFB₁ exposure in Fischer 344 rats, and to assess exposures to AFB₁ in different human populations based on the measurement of serum AFB₁-lysine adduct.

The hypothesis of the dissertation

We propose to establish a global network of collaborators to study aflatoxin exposure using serum AFB₁-lysine adduct as a biomarker. Our **working hypothesis** is that levels of AFB₁-lysine adduct in human serum are highly stable and correlated with aflatoxin intake, and will be a reliable biological response indicator for human aflatoxicosis in high-risk populations.

The **immediate goal** of this dissertation project is to develop a highly sensitive, non-antibody, non-radioactive, and non-mass spectrometry (MS) based analytical method for rapidly measuring serum AFB₁-lysine adduct. The method will be validated in animals treated with single or repeated doses of AFB₁ and in archived blood samples previously collected from high risk populations in developing countries. Analysis will be performed in samples from human biomonitoring and/or intervention studies conducted

in China, Ghana, and Burkina Faso. Our **long-term goal** is to establish a global network for studying aflatoxins exposure and evaluating the efficacy of chemoprevention strategies based on measuring the molecular biomarker of AFB₁-lysine adduct in serum.

The specific aims of this dissertation include:

Phase I: To develop a highly sensitive, non-antibody, non-radioactive, and non-mass spectrometry (MS) based HPLC-fluorescence method for rapidly measuring serum AFB₁-lysine adduct;

Phase II: To validate the method in animals (rats) treated with a single dose (50, 250, and 1000 µg/kg body weight) or repeated doses (5, 10, 25, and 75 µg/kg body weight for 5 weeks) of AFB₁ and in archived human blood samples previously collected from high risk populations in US, Ghana, and China.

Phase III: To evaluate the integrative toxicological approach in AFB₁ treated rats. The adverse effects of AFB₁ on the splenic immune functions, liver preneoplastic changes, and serum biochemical alternations will be compared to concurrent measurement of serum AFB-Lys adduct levels.

Phase IV: To assess human exposure in samples from human biomonitoring and intervention studies conducted in China, Malaysia, the United States, Ghana, Burkina Faso, Uganda and Haiti.

This dissertation is intended to validate an integrative toxicological evaluation of AFB₁ exposure in Fischer 344 rats and the assessment of AFB₁ exposure in different human populations using the biomarker serum AFB₁-lysine adduct. AFB₁ exposure has been linked to many human diseases including aflatoxicosis, Reye syndrome, hepatocellular carcinoma, immune and growth retardation in children. The role of AFB₁

biomarkers including serum AFB₁-lysine in the process of clinical diseases or adverse health effects related to AFB₁ remains largely unknown. The effects of AFB₁ exposure on the histological, immunohistological, clinical biochemical parameters, as well as on the cell-specific cytokine secretion in splenic lymphocytes will be explored in Fischer 344 rats through both single-dose and repeated-dose treatment protocols. Temporal correlations among these changes will be explored. The temporal changes in serum AFB₁-lysine adduct will be monitored and the kinetic parameters (e.g., peak time, half-life and elimination kinetics) of this biomarker will be validated using the non-antibody method. The results from the animal experiment will increase our understanding of the potential use of molecular biomarker into the toxicity assessment of AFB₁. Such data could be useful for delineation of functional and microstructural changes in liver and for evaluation of various intervention strategies to diminish the acute and chronic toxic effects due to AFB₁ exposure. Human exposure data from populations worldwide can serve as the first step towards a global network that reflects current human AFB₁ exposure. Such data are also important for identifying high risk populations and can be used, together with the human demographic information, to reveal potential correlations with other human disease or adverse health effects.

CHAPTER 2

LITERATURE REVIEW

General Overview

Aflatoxins were discovered in the early 1960s as the causative agent of turkey X disease in Great Britain, which caused the death of thousands of young turkeys and ducklings that were fed with a contaminated peanut meal originating in Brazil^{11, 12}. The most significant changes of turkey X disease, as described by Wannop¹³, were in the liver, including areas of eosinophilic parenchymatous cells with homogenous cytoplasm, swollen parenchymatous cells and their nuclei, obliteration of many sinusoids, loss of the structure of the liver cord, biliary proliferation, necrosis and regeneration near the portal tract and large vacuoles. In the contaminated feed, the fungi *Aspergillus flavus* and *Aspergillus parasiticus* were identified as the culprits that produce toxins^{14, 15}. The toxins were characterized chemically and were designated aflatoxins¹⁶. From a retrospective standpoint of view, the outbreaks of turkey X disease were preceded by a number of less well-described episodes of epizootics in different animal species¹⁷. The outbreak of turkey X disease marked a new era for the intensive research on the toxicology and biochemistry of mycotoxins.

The aflatoxins species B and G are designated according to their fluorescence, e.g. B species (B₁, B₂) show blue fluorescence and G species (G₁ and G₂) show greenish-yellow fluorescence under UV light. Chemical synthesis demonstrated that aflatoxins are a highly substituted coumarin structure containing a dihydrofurofuran moiety (Figure

2.1). The structural characterization and total synthesis of aflatoxins were accomplished in 1963¹⁶, which greatly enhanced mechanistic research of their toxicity and carcinogenicity.

Extensive assays were conducted in a wide range of animal models to assess the acute and chronic toxicity of aflatoxins starting from the early 1960s^{18, 19}. Among the naturally occurring aflatoxins, aflatoxin B₁ (AFB₁) is the most potent in terms of acute toxicity, mutagenicity, and carcinogenicity¹. Acute exposure to a single large dose of AFB₁ leads to massive liver damage, including edema, periportal necrosis and bile duct proliferation, and death in rodents. Along with the histological changes, altered liver enzyme activities (ALT, AST) and induction of altered foci of hepatocytes (preneoplastic lesion) are frequent outcomes within a few days to several weeks after administration^{20, 21}. Chronic studies demonstrated that aflatoxins are complete carcinogens and that increasing the dose decreases the latency for tumor development²². An early study showed that the effective carcinogenic dose of AFB₁ is less than 0.5 mg per rat²³. Repeated low dose exposure is more efficient than a single large exposure, which is consistent with the property of a complete carcinogen. A metabolism study led to identification of the major AFB₁-DNA adducts in 1977²⁴. Dose-dependent AFB₁-DNA adduct formation in rat liver occurs over a range of doses up to several orders of magnitude. The early development of analytical methods for quantifying aflatoxins from foods represents extensive international collaborations. These physical, chemical, and immunologic analytical methods¹⁰ greatly facilitated the regulation, monitoring, and reducing of aflatoxins contamination in foods. Based on these methodologies, many epidemiologic studies have been conducted to evaluate the association of aflatoxin intake

and the development of hepatocellular carcinoma (HCC) in human populations⁴. The development and validation of aflatoxin biomarkers, including AFB₁-DNA adduct, serum aflatoxin-albumin adduct, aflatoxin M₁, and aflatoxin B₁ mercapturic acid, have provided the tools to evaluate the molecular epidemiology of aflatoxin exposure and risk to individuals in China and sub-Saharan Africa²⁵. These lines of investigation are still being conducted in current efforts to develop effective intervention strategies to mitigate the health impacts of aflatoxin exposures. The human data from such studies and accumulated experimental evidence provided the basis for the classification of AFB₁ and its mixtures as a human carcinogen by the International Agency for Research on Cancer (IARC) in 1993¹. The time line for key events in the discovery, toxicological evaluation, molecular epidemiology, and regulation of aflatoxins has been nicely shown in a recent review (Figure 2.2)⁴.

Aflatoxin exposure remains a largely ignored global issue. It is estimated that approximately 4.6-28.2% of global annual HCC cases are attributed to aflatoxin exposure alone³. In the developing world, more than 40% of all disease burden are estimated to be associated with aflatoxin exposure². These chronic health effects together with the growth impairment in children⁵ point to the urgent needs for effective intervention strategies to reduce the risks associated with aflatoxins. After the severe outbreak of aflatoxicosis in Kenya²⁶, an international workshop has initiated public health strategies for reducing aflatoxins exposure in the developing countries²⁷. Intervention strategies, which aim to block, retard, reverse the exposure and risk of aflatoxins, have been a focus of research that is ongoing. The approaches of preventions that have been studied include improving of postharvest storage conditions, using trapping agent (e.g., clay) to reduce

bioavailability or agents (e.g., green tea polyphenol, chlorophyllin, and dithiolethiones) to alter the metabolic pathway to reduce risk caused by aflatoxins exposure^{2, 28, 29}.

The strategy to define the risks from aflatoxins exposure and to design approaches to prevent their effects serves as a template for the development, validation and application of molecular biomarkers for other environmental agents or their resulting chronic diseases. This strategy also provides a model for the design of future studies for risk assessment of exposure to other environmental agents. Key elements of such a strategy has been summarized by Kensler et al.⁴, which include: the use of animal models to mimic the human disease; designing observational studies associating exposure with disease; development of mechanism-based molecular biomarkers in animal models; validation of these biomarkers in animals by dose-response studies; validation of transitional human studies; and association of these biomarkers with disease outcome in prospective studies.

Toxicokinetics of Aflatoxin B₁

Absorption

The rate of absorption of AFB₁ may be affected substantially by the route of exposure. Being a low molecular, lipophilic molecule, the absorption of AFB₁ from the gastrointestinal (GI) tract is fast and efficient. The distribution of radioactivity after oral or intraperitoneal (ip) injection of [¹⁴C] AFB₁ in Male F344 rats is similar³⁰, suggesting efficient absorption after ingestion. Both species and strain differences play a role in the absorption of AFB₁ through oral administration. For example, the rate of stomach emptying in rat is slower than that in mouse³¹, which may affect the rate of absorption through the GI tract. Within 24 h of an oral administration of [¹⁴C]AFB₁, up to 20 to 60%

of the radioactivity was eliminated in feces in F344 rats ³⁰, whereas approximately 10-30% of total radioactivity was excreted in the bile in Wistar rats ³².

A previous study showed that the rate of AFB₁ uptake in intestinal tissue is proportional to AFB₁ concentration ³³, suggesting a mode of passive diffusion; intestinal absorption of aflatoxin is more efficient in suckling than older rats. Therefore, lipophilicity of aflatoxins and the composition of the intestinal epithelium may be important determinants of aflatoxin absorption. The aflatoxin metabolism that occurs in the GI mucosa represents a major AFB₁ detoxification site as approximately 20-25% of the ingested dose can reach the liver ³⁴. This is supported by a report that cytochrome P450 (CYP) 3A4 protein content is 3 times higher in the enterocyte homogenates than in the liver homogenates ³⁵. Reabsorption of AFB₁ conjugates through enterohepatic recirculation was found especially at higher doses (0.5-2.5 mg/kg) but not at low dose (0.05 mg/kg) ³². A study of AFB₁ absorption in lactating Holstein dairy cows showed that after a single bolus of [³H]AFB₁ (4.89 mg), plasma AFM₁ can be detectable at 5 min (10.4 ng/L) after treatment, which peaked at 25 min (136.3 ng/L) after treatment ³⁶. The author suggested that AFB₁ uptake and metabolism may occur in mouth or oesophageal mucous membranes, before the rumen compartment. This study also demonstrated that vaginal implant of AFB₁ lead to plasma detectable AFB₁ and AFM₁ at 30 min after treatment, suggesting an absorption mediated by passive diffusion.

Human exposures to aflatoxins occur mainly through ingestion of contaminated foods, including corn, peanuts, cottonseed, tree nuts and pepper. This issue is most serious in developing countries that lack proper processing procedures, storage facilities, and food-safety monitoring ^{37, 38}. Infants or young children can be exposed to AFM₁

through ingestion of breast milk or dairy products that are contaminated by AFB₁. Occasionally, the meat products intended for human consumption can be contaminated by aflatoxins, which represent a much less common source of exposure³⁹.

Although dietary exposure to AFB₁ has been widely recognized, evidence has been accumulated to suggest potentially high risks of occupational exposure to AFB₁ through inhalation⁴⁰. During the processing of foods such as rice or maize, workers are exposed to aflatoxin producing fungi through dust where aflatoxins have been isolated⁴¹,⁴². When human sera were collected, the AFB₁-albumin adduct levels were detected to be higher in workers exposed to dust with detectable aflatoxins than workers who were not exposed⁴³. It was reported that AFB₁ was present in the lungs of one textile and two agricultural workers who died from pulmonary interstitial fibrosis⁴⁴, and that AFB₁ was found in the lungs of chemical engineers who had worked for 3 months on a method for sterilizing Brazilian peanut meal contaminated with *Aspergillus flavus* and died of alveolar cell carcinoma⁴⁵. Thus, inhalation is an important route for occupational exposure. Recently, airborne aflatoxins have been assessed for their levels during a grinding phase. When a batch of imported shelled peanuts imported from Vietnam was ground wet, the level of aflatoxins was shown to be negligible (0.11 pg/m³)⁴⁶. However, the contamination level in the peanuts or shells was not provided, which may compromise this conclusion.

Human data with regards to the rate of absorption of aflatoxins are not as abundant as that in animal species. Whether AFB₁ can penetrate isolated human epidermis (stratum corneum plus viable epidermis) was tested in vitro⁴⁷. [¹⁴C] AFB₁ (7.5 - 9.3 µg) was applied to the stratum corneum of epidermal disks mounted in Teflon

diffusion cells. [¹⁴C]AFB₁ penetrated through the isolated epidermis in chemically unaltered form, at a slow rate under nonoccluded conditions but at a rate that was approximately 40 times greater when the skin was occluded. The maximum velocity of penetration was 0.63 ± 0.71 and 27.31 ± 10.15 pmol/h under conditions of exposure to ambient conditions and occlusion, respectively. Chloroform extractable radioactivity accounted for $82.5 \pm 3.7\%$ of the total penetrating radioactivity in the receptor fluid of the diffusion cells. More than 95% of the applied dose penetrated epidermis within 46 h. Total recovery was $98.6 \pm 6.4\%$, as estimated by a percentage of the applied radioactivity. These results demonstrate the features of AFB₁ absorption through skin: slow and near complete, but fast under conditions of occlusion which increase the hydration of the skin. Human volunteers that were orally give low doses of [¹⁴C] AFB₁ (30 ng, 5 nCi) demonstrated a peak total radioactivity in plasma at 1 h after exposure⁴⁸, suggesting rapid absorption through the route of gastrointestinal (GI) tract.

Distribution

Different animal models had been used to compare the biological fate of free AFB₁ in blood. A single iv dose of [¹⁴C]AFB₁ at 0.3, 1 or 1.5 mg/kg was given to the rhesus monkey, Sprague-Dawley rat and Swiss-Webster mouse, respectively, which are approximately 1/10 of the LD₅₀ doses, respectively. The apparent volumes of distribution were calculated to be 114%, 47%, and 28% for the monkey, rat and mouse, respectively⁴⁹. This ranking is consistent with the susceptibility of these species to AFB₁ toxicity. Liver is the major site for concentration of AFB₁, especially at low doses. The radioactivity in liver was 5-15 fold higher than other tissues at 30 min after administration of [¹⁴C]AFB₁³⁰. Within a few hours, most of the AFB₁ in liver is bounded

to macromolecules. For instance, it was found that about 15% of the AFB₁ dose is retained in rat liver at 2 h after treatment, and 70% of the retained dose is covalently bound adducts⁵⁰.

The tissue distribution of AFB₁ appears to be species-specific. After 2 weeks administration of [¹⁴C]AFB₁ to broiler chickens and 5 h after the last dose, about 96.64% of the ¹⁴C was excreted. Of the ¹⁴C retained, 11.04, 9.83, 4.30, 12.52, 31.66, and 30.63% were detected in the blood, liver, heart, gizzard, breast, and leg, respectively⁵¹. In a 35 days feeding trial with AFB₁ in swine, the highest concentrations of aflatoxins were found in kidney and liver, and substantially lower concentrations were found in muscle and adipose tissues⁵². In rats that were given ip doses of [¹⁴C] AFB₁, radioactivity was present at maximum levels in liver and kidney 0.5 h after administration, with small amounts present in other organs. The concentration of radioactivity was 5 to 15 times greater in liver than in other tissues, and at the end of 24 h, the liver contained an amount equal to the content of the remainder of the carcass³⁰. This finding is associated with the relative tissue specificity of AFB₁ as a hepatotoxin in rats.

Accumulated experimental evidence has shown that aflatoxin or its metabolites can reach the developing fetuses. In mice that were treated with [¹⁴C]AFB₁, radioactivity were found in the eyes of the fetuses although their livers did not accumulate radioactivity⁵³. Similar results were found in pregnant mice, the fetuses showed distinct uptake of radioactivity in the pigment layer of the eyes and in the nasal mucosa, the liver had only small amounts of radioactivity although the liver in the mother had a very high concentration⁵⁴. Transplacental transfer of aflatoxin in humans have been confirmed by many studies showing the presence of aflatoxin and/or its metabolites in the cord sera⁵⁵⁻

⁵⁸. Although several metabolites have been identified in the cord sera, it is not clear whether they are formed in placenta or are of maternal origin. A recent study has reported that aflatoxicol is the only metabolite detected in both human placenta perfusion and *in vitro* incubation studies ⁵⁷. If aflatoxicol is in fact the only metabolite in placenta, then aflatoxin metabolites found in cord sera (e.g., aflatoxin-albumin adducts) may come from the maternal blood rather than being a product of placenta metabolism. Although less mutagenic than AFB₁, aflatoxicol still has higher mutagenicity than other metabolites ⁵⁹, therefore, the placenta may not be considered as a significant protective barrier for fetal aflatoxin exposure.

Biotransformation

The intestinal mucosa serves as a primary site of metabolism, reducing the amount of aflatoxin available to the liver. Once absorbed, the liver is the major site for biotransformation and metabolism of aflatoxins. The aflatoxins undergo phase I and phase II metabolic process in liver. Phase I reactions include hydroxylation, epoxidation and demethylation of aflatoxins, producing aflatoxin M₁, Q₁, P₁. Different members (1A2, 3A4, 3A5 and 3A7) of the cytochrome P450 (CYP450) enzyme family have been found to be responsible for aflatoxin metabolism ⁶⁰. Compared with CYP 3A4, human liver microsomal CYP 1A2 is reported to have a higher affinity toward low AFB₁ concentrations that are more reflective of dietary exposures ⁶¹. This differential metabolism of these two isoforms was also found in human bronchial cells ⁶². On the other hand, a significant correlation between CYP 3A4 expression (but not CYP 1A2) and AFB₁-DNA adduct levels was found in adult human livers ⁶³. Using a hepatic abundance model that takes into account the specific kinetic parameters and the

expression levels of these CYP450s, Kamdem et al. ⁶⁰ compared the relative production rates of the carcinogenic metabolite, AFB₁-8,9-epoxide, by different CYP450s. They found that CYP 3A4 is the most important determinant towards the epoxide formation, followed by CYP 3A5, 3A7, and 1A2 ⁶⁰. In addition, CYP450 specific metabolites have been found, for instance, 3A4 catalyzes hydroxylation of AFB₁ to produce AFQ₁ while 1A2 generates another hydroxylated metabolite, AFM₁. Taken together, CYP1A2 has a higher affinity but a lower capacity towards metabolic activation of AFB₁ and conversely CYP3A4, has a much larger capacity but a lower affinity. Unlike the reduced metabolite aflatoxicol that is excreted in urine without further metabolism or in feces after glucuronyl conjugation ⁶⁴, AFM₁ can be further metabolized to form AFM₁-8,9-epoxide. This reactive epoxide binds to DNA producing AFM₁-N⁷-guanine, which can be detected in the liver DNA and urine of tree shrew and rats 24 h after treatment with AFB₁ (400 µg/kg) ⁶⁵. Tree shrews produced twice as much AFM₁-N⁷-guanine adducts compared to rats (0.74 ± 0.14 pmol/mg DNA vs 0.37 ± 0.07 pmol/mg DNA). However, in the rat liver, the ratio of AFB₁-DNA adducts to AFM₁-DNA adducts is about 20 (6.56/0.42 pmol/mg DNA). This metabolic difference might account for the sensitivity of rat to AFB₁ hepatocarcinogenesis.

The lung is also a target organ for aflatoxins. Tissues derived from various portions of the lung have been used as *in vitro* models to study AFB₁ metabolism and activation ⁶⁶. AFB₁ can be metabolically activated by intact human tracheal epithelium from the bronchus and in human epithelioid lung cells *in vitro*. The two major AFB₁-DNA adducts are also formed in the pulmonary tract in ratios similar to those seen in

hepatic tissues of AFB₁-treated animals⁶⁶. Inhalation of AFB₁ by rats for 20-120 min resulted in an increase in AFB₁-N⁷-Gua adduct in a dose-dependent manner.

Of the metabolic products produced during phase I reactions, the most biologically important is the highly reactive AFB₁-8,9-epoxide⁶⁷. This metabolite has been found to exist in 2 stereoisomers, of which the *exo* isomer is much less stable in water and approximately 500 times as mutagenic as the *endo* isomer. The increased mutagenicity of the *exo* isomer is attributed to its steric configuration that allows covalent binding of DNA, almost exclusively between the N⁷ of guanine and the C⁸ of the 8,9-*exo*-epoxide⁶⁸⁻⁷¹. The rate of conjugation of glutathione (GSH) with AFB₁ *exo*-epoxide is an important factor in determining the species variation in risk to aflatoxin hepatocarcinogenicity. The *endo*-epoxide was found to be a good substrate for GSH conjugate formation in rat liver cytosol while mouse liver cytosol conjugated the *exo*-epoxide almost exclusively⁷². This difference may partly account for the resistance of mouse to the hepatocarcinogenicity of AFB₁.

The Phase II biotransformation leads to conjugation of phase I metabolites of aflatoxins forming less toxic or carcinogenic products than the parent compound. This reaction also increases the polarity and facilitates the excretion of the conjugates from urine and feces. The detoxification pathway of aflatoxin is shown in Figure 2.3⁷³. Liver cytosolic glutathione S transferase (GST) plays a critical role in the detoxification of AFB₁-8,9-epoxide by conjugating glutathione with the epoxide⁷⁴. Glutathione conjugation is an important reaction that determines the susceptibility of different species to the toxicity of AFB₁, with a most striking difference observed between the mouse and the rat⁷⁵.

Genetic polymorphism of the metabolic enzymes also affects the susceptibility to the toxicity and carcinogenicity of aflatoxins in different animal species or possibly ethnic human groups ⁷⁶. CYP3A5, GST, and epoxide hydrolase (EH) polymorphisms are of biological significance in particular. CYP3A5 exhibits genetic polymorphism known as CYP3A5*3, CYP3A5*6, and CYP3A5*7 ^{77, 78}. It is estimated that approximately 90% Caucasians, 67% Japanese, and 30-45% African-Americans are CYP3A5 low expressors ⁷⁷⁻⁷⁹. Consistently, Wojnowski et al. reported increased plasma levels of AFB₁-albumin adducts in CYP3A5 high expressors from Gambia, particularly in those with low concomitant CYP3A4 expression ⁸⁰. It is conceivable that the individual expression levels of CYP450s producing AFB₁-8,9-epoxide could affect the individual risk to develop HCC.

Detoxification pathways prevent formation of DNA adducts, which involve the glutathione conjugation of reactive metabolites by GST or hydrolysis by microsomal epoxide hydrolase (mEH) ^{81, 82}. There are two types of genetic polymorphic GSTs that are responsible for AFB₁ detoxification: GST- μ encoded by the *GSTM1* gene and GST- θ encoded by *GSTT1*. Homozygous deletion of part of these genes (null genotype) results in enzyme deficiency and might therefore lead to hampered detoxification ⁸³. The frequencies of homozygous *GSTM1* and *GSTT1* deletion carriers are very high (i.e., 20-50%) in most populations studied to date ⁷⁸, and the risk conferred to individuals who carry homozygous deletions in *GSTM1* and *GSTT1* is increased when the interactions of these genotypes with other environmental carcinogens are involved (e.g., odds ratio increasing from less than 2 to 3-5) ⁸⁴. In several studies, it has been shown that only the *GSTM1* genotype carriers are at increased risk of HCC in populations exposed to

aflatoxins^{76, 85-87}, although GST- θ has also shown high efficiency for conjugation of glutathione to AFB₁-8,9-epoxide^{88, 89}. It has been noted that the deletion genotypes are so common in the general population that the proportion of cancer attributable to these variants may be large, however, these genotypes may be less suited for individual cancer risk assessment because of their relatively small contribution to the absolute risk of cancer⁸⁴. Besides carcinogen metabolizing enzymes, the polymorphism of DNA repair (*XRCCI*) enzyme has been shown to be related to the risk of HCC development in Gambia, West Africa⁹⁰ where aflatoxin exposure is high. In that hospital based case-control study (216 HCC cases and 408 controls)⁹⁰, a multivariable analysis (adjusting for demographic factors, hepatitis B virus, hepatitis C virus, and TP53 status) revealed that the heterozygote *XRCCI*-399 AG genotype were significantly associated with HCC (OR, 3.18; 95% CI, 1.35-7.51), although the prevalence of variant genotypes was generally low.

Human mEH catalyzes the conversion of a broad array of reactive epoxide metabolites to more polar trans-dihydrodiol derivatives. The gene has been shown to exhibit polymorphism, including variation in the coding region leading to amino acid substitutions at positions 113 (Y/H) and 139 (H/R)⁹¹. *In vitro* studies have shown that human mEH provides no statistically significant changes to the rate of hydrolysis of AFB₁, benzo[a]pyrene-4, 5-oxide, and cis-stilbene oxide^{92, 93}, compared to the rate of spontaneous hydrolysis or the variants of mEH proteins. Such evidence suggests that human mEH may not play an important role in detoxification of AFB₁.

Excretion

Previous studies showed that after an oral or iv dose of [³H]AFB₁ in Sprague-Dawley rats, more than half of the total radioactivity is excreted into feces⁹⁴. The major biliary metabolite was AFB₁-glutathione, accounting for 49-57% of the total biliary excretion, and AFP₁-glutathione accounts for 4-15% of total biliary radioactivity in Sprague-Dawley rats⁵⁰. Urinary excretion represents another important route of excretion. Approximately 10-20% of a single dose of AFB₁ is excreted in urine 24 h after ip administration to the rat, and the major urinary metabolites are determined to be AFM₁, AFP₁, and AFB₁-N⁷-guanine⁹⁵.

Using a monoclonal antibody specific for aflatoxins, Groopman et al. detected the urinary metabolites of AFB₁ in urine from AFB₁ treated rats and human subjects from Guangxi province of China where aflatoxin exposure is high⁹⁶, the major urinary metabolites are AFB₁-N⁷-guanine, AFM₁ and AFP₁ in both rats and humans. The fecal and urinary excretion of 3 AFB₁ metabolites, AFM₁, AFQ₁, and AFB₁-N⁷-guanine in humans has been assessed in another study⁹⁷. Eighty-three young Chinese males were selected from 300 people based on detectable urinary AFM₁. The fecal excretion of AFQ₁ (median, 137 ng/g fresh weight) was approximately 60 times higher than that of AFM₁ (median, 2.3 ng/g). In urine, the median levels of AFQ₁, AFM₁, and AFB₁-N⁷-guanine were 10.4, 0.04, and 0.38 ng/mL, respectively. A subgroup (n = 14) with HBV infection had significantly higher fecal concentrations of AFQ₁ (*P* = 0.043) and AFM₁ (*P* = 0.001) than those who were not HBV positive. That study demonstrated that AFQ₁ is excreted in urine and feces at higher levels than AFM₁, and feces are an important route of excretion

of these AFB₁ metabolites. However, whether AFQ₁ should be suitable to be used as a marker for exposure and risk of dietary aflatoxin intake remains unclear.

The nutrient status seems to affect the excretion of aflatoxins. Children with kwashiorkor (n = 5) and marasmic kwashiorkor (n = 7) were investigated for the urinary and fecal excretion of aflatoxins⁹⁸. Aflatoxin-free diet consisting of maize meal and milk powder was provided, and urine and feces samples were collected on admission and day 4 and 10. The children with kwashiorkor excreted aflatoxins in stools for up to 9 days and ceased urinary excretion after 2 days. The children with marasmic kwashiorkor excreted aflatoxins in stools for 6 days and ceased urinary excretion after 4 days. In stools, AFB₁ was most frequently detected in kwashiorkor and least frequently in marasmic kwashiorkor.

Toxicity of Aflatoxin B₁

Acute Toxicity

The aflatoxins were discovered because of their acute toxicity in poultry. The acute toxicities of aflatoxins and their metabolites have been tested in different animal species. Overall, AFB₁ is the most toxic form among the naturally occurring aflatoxins, as determined in duck^{99, 100} and rat⁹⁹. The acute toxicity varies up to several orders with regards to the median lethal dose (LD₅₀), as shown in Table 2.1^{99, 101-104}. It can be seen from Table 2.1 that age is an important factor affecting the acute toxicity to AFB₁: young mouse and rat are more susceptible to the toxicity of acute AFB₁ exposure than their adults. The rank of potency in terms of acute toxicity from high to low is B₁ > G₁ > B₂ > G₂ in rat and duck⁹⁹. Acute exposure to AFB₁ causes a series of symptoms, including abdominal pain, nausea, jaundice, bleeding in the GI tract, and even death (aflatoxicosis).

The histopathological effects and biochemical changes induced by AFB₁ exposure have been extensively studied in animal models^{18, 19, 105}. However, only a few studies^{20, 106} reported the sequential biochemical and histological changes, in which rats were given either one single large dose of 3 mg/kg body weight (BW) or 7 mg/kg BW. In particular, no studies have been done to explore the correlations between biochemical and histopathological alterations with specific molecular targets via single or repeated treatment protocols. Rat liver GST-P positive (GST-P⁺) foci are a specific and reliable preneoplastic marker used widely in animal carcinogenesis models¹⁰⁷. Short-term exposure to AFB₁ induces GST-P⁺ hepatocytes or foci in rat liver both independently and synergistically with other toxins, such as fumonisin¹⁰⁸ and microcystin LR¹⁰⁹. However, the sequential development of GST-P⁺ foci caused by AFB₁ as well as the correlations with biochemical and histological changes in animal models remains largely unknown.

Human Aflatoxicoses

Historically, the outbreaks of human aflatoxicoses have been reported especially in the developing countries^{2, 110}. These outbreaks have been frequently reported from India^{111, 112}, Kenya¹¹³, and Malaysia^{114, 115}. A case was also reported in Uganda¹¹⁶. These acute aflatoxicoses are as summarized in Table 2.2. During the first reported outbreak in India (1975), analysis of contaminated samples showed that affected people could have consumed between 2 and 6 mg of aflatoxin daily over a period of one month. Affected people (n = 397) showed jaundice, rapidly developing ascites, portal hypertension, and a high mortality rate. This outbreak was associated with consumption of heavily contaminated maize¹¹¹. The second outbreak in India involved 994 cases who had consumed aflatoxins contaminated maize¹¹². Willis et al. reported a case that a 25-

year-old woman who attempted suicide twice by ingesting mixtures containing 15-45% AFB₁. She ate 5.5 mg over 2 days and 6 months later 35 mg more over 2 weeks. Following the first exposure, she was admitted to hospital with a transient, nonpruritic, macular rash, nausea and headache; the second time she reported nausea only ¹¹⁷. On both occasions, physical and laboratory examinations were normal and liver biopsies appeared normal by light microscopy. After 14 years follow-up examination, the women remained healthy and no any signs or symptoms of disease or lesions were revealed. These findings suggest that the hepatotoxicity of AFB₁ may be lower in well nourished persons than in experimental animals or a longer latent period than 14 years would be expected for tumor formation. The outbreak of aflatoxicosis in Kenya (2004 and 2005) represented the most severe case of aflatoxicosis that caused 125 deaths and 317 cases which resulted from the ingestion of heavily contaminated maize and maize product ¹¹⁸.

Reye Syndrome

Reye syndrome is characterized by sudden brain damage and liver function problems of unknown cause. Although the syndrome has occurred in children who have been given aspirin or phenothiazines when they have chicken pox or the flu ¹¹⁹, aflatoxins have also been suggested as a potential etiological factor in Reye syndrome because of the similar changes induced by aflatoxins, such as encephalopathy and fatty degeneration of viscera ¹²⁰. Concurrent aflatoxin exposure and Reye syndrome in children were reported in children in Czechoslovakia. Of the 26 children with Reye syndrome, all of them had detectable AFB₁ in liver and 15 children also possessed detectable AFM₁ ¹²¹. None of the 25 control children had detectable AFB₁ or AFM₁ in the liver ¹²². It was also reported in New Zealand that two children with Reye syndrome had detectable AFB₁ in

the liver tissues ¹²³. Several reports in the United States showed that AFB₁ is 100% detectable in blood or liver tissues of children with Reye syndrome ^{124, 125}. In Thailand, a case-control study showed that AFB₁ was more frequently detected in the liver, brain, and kidney tissues from children who died of Reye syndrome than the control children who died of other diseases ¹²⁶. An association between AFB₁ exposure and Reye syndrome is suggested by these case reports and studies. However, since AFB₁ is not regularly recoverable from cases of Reye syndrome at a high rate, the causal effect was questioned ¹²⁷. Therefore, the causal effect of AFB₁ remains to be confirmed in more carefully designed studies particularly with a prospective design.

Chronic Toxicity: Hepatocellular Carcinoma

Based on sufficient evidence from animal experiments and human epidemiological studies, aflatoxins have been established as a known human carcinogen ¹. Approximately 4.5 billion people worldwide are estimated to be chronically exposed to large amounts of aflatoxins ². Among the 550,000-600,000 new HCC cases worldwide each year, about 25,200-155,000 may be attributable to aflatoxin exposure, as estimated by Liu and Wu ³. There is a broad ecologic association between areas of high aflatoxin exposure (such as China, Southeast Asia, and parts of Africa) and HBV, which is endemic in these areas; rates of HBV positivity in Gambia, China, and Guinea are reported to be 15% ¹²⁸, 14–20% ¹²⁹, and 10% ¹³⁰, respectively. Independently, each factor significantly increases the relative risk of cancer, and most studies report them to be synergistic ¹³¹, increasing the risk of cancer 25 to 30-fold; there is, however, one epidemiologic study that contradicts those findings ¹³². Aflatoxin exposure multiplicatively increases the risk for development of human HCC for people chronically

infected with hepatitis B virus, as demonstrated by several human studies that employed molecular biomarkers (e.g., urinary AFB₁ metabolites and serum aflatoxin-albumin adducts or codon 249 mutations) to improve individual risk assessment¹³³⁻¹³⁶. Recently, a systematic review and meta-analysis reported the population attributable risk (PAR) of aflatoxin-related HCC¹³⁷. The PAR of aflatoxin-related HCC was estimated to be 17% overall, which increased to 23% when the one study that contributed most to heterogeneity in the analysis is excluded. The summarised odds ratios (OR) of HCC with 95% CIs are 73.0 (36.0-148.3) from the combined effects of aflatoxin and HBV, 11.3 (6.75-18.9) from HBV only, and 6.37 (3.74-10.86) from aflatoxin only. These data indicate that aflatoxin multiplicatively interacts with HBV to induce HCC and reducing aflatoxin exposure to non-detectable levels could reduce HCC cases in high-risk areas by about 23%. Whereas the importance of this synergy to cancer is well recognized^{138, 139}, its importance to other health issues is largely unknown and clearly should be established.

A number of possible mechanisms for the interaction have been suggested¹⁴⁰⁻¹⁴², which include, among others:

- Chronic HBV infection may induce the CYP450s that metabolize inactive AFB₁ to the mutagenic AFB₁-8,9-epoxide.
- Nuclear excision repair, which is normally responsible for removing AFB₁-DNA adducts, is inhibited by HBVx protein.
- HBVx protein may also increase the overall frequency of DNA mutations, including the *p53* 249^{ser} mutation.
- The fixation of AFB₁-induced mutations in the presence of liver regeneration and hyperplasia induced by chronic HBV infection.

- The predisposition of HBV-infected hepatocytes to aflatoxin-induced DNA damage.
- An increase in susceptibility to chronic HBV infection in aflatoxin-exposed individuals.
- Oxidative stress exacerbated by co-exposure to aflatoxins and chronic hepatitis infection.

These mentioned mechanisms remain to be confirmed and inconsistent results have been found, e.g., an *in vitro* study has recently showed that in HepaRG cells, AFB₁ exposure (up to 5 µM) decreases HBV replication, whereas AFB₁ induced DNA damage (AFB₁-DNA adduct formation) and subsequent *p53* alteration (mutation at codon 249) is not affected by the presence of the virus¹⁴³.

Immune Toxicity

Immune suppression in experimental animals has been a common outcome of aflatoxin exposure. AFB₁ causes immune suppression in diverse laboratory and domestic animal models, with the cell-mediated immunity being the most affected¹⁴⁴. A considerable amount of evidence from both *in vivo* and *in vitro* studies has demonstrated that AFB₁ affects immune functions^{144, 145}. AFB₁ decreases splenic helper T cells and antibody response to sheep red blood cells (SRBC) in C57BL/6 mice¹⁴⁶ and inhibits production of IL-2, IL-3, and IFN-γ by splenic macrophages¹⁴⁷. Phagocytosis and production of nitric oxide and TNF-α by macrophages are inhibited by AFB₁ both *in vivo* and *in vitro*^{148, 149}. In weaning rats, exposure to AFB₁ decreases spleen and thymus weights, inhibits phagocytic capacity of macrophages¹⁵⁰ and suppresses delayed type hypersensitivity response¹⁵¹. Dietary exposure of rats to AFB₁ (0.01-1.6 ppm) produces

inhibitory effects on splenic T and B lymphocytes as well as IL-1, IL-2, and IL-6 production by splenocytes¹⁵². In addition, chick and swine are also susceptible to AFB₁ induced alterations in cell-mediated immune response¹⁵³⁻¹⁵⁵ and AFB₁ has been shown to stimulate the expression of pro-inflammatory cytokines such as TNF- α , IFN- γ , and IL-6¹⁵⁶. Besides dietary exposure, inhalation of aflatoxins is also capable of reducing suppression of pulmonary and systemic host defense in mice and rats¹⁵⁷.

The complement components, as part of the innate immune system, helps antibodies and phagocytic cells to clear pathogens from an organism. The activities of complements have been shown to be depressed due to aflatoxin exposure in various animal models¹⁵⁸⁻¹⁶¹, including chicken, Guinea pig, swine and cattle. The observed depressive effects on complement activity may be related to the ability of aflatoxins to interfere with protein synthesis. Early research had also shown that aflatoxin exposure reduces the resistance of several animal species to bacterial, viral, fungal, and parasitic infection^{162, 163}. Noteworthy, beneficial effects of aflatoxin exposure to bacterial infection were also reported. A nonspecific humoral factor appeared to be affected in the aflatoxin-impaired acquired immunity to *P. multocida* in turkey poults because serum from normal or immunized turkeys was beneficial in overcoming the impairment¹⁶⁴. Similarly, oral exposure to aflatoxin decreased bacterial counts in intestinal tract and mesenteric lymph nodes of hamsters that were infected with mycobacterium paratuberculosis, although aflatoxin treated hamsters showed inhibitory growth and liver lesions¹⁶⁵. The effects of aflatoxins on the resistance of animals to infectious disease agents are somewhat variable, depending on the organism, toxin dose and formulation, species, and perhaps sensitivity of the test¹⁶⁴.

Only circumstantial evidence supports the causative effects of AFB₁ exposure to suppression of human immune functions. Of significance is the cohort study investigating the effects of aflatoxin exposure on the immune functions in Gambian children ¹⁶⁶. The level of secretory IgA (sIgA) in saliva was significantly lower in children with detectable levels of serum AFB₁ albumin adduct than those with non-detectable levels. Jiang et al. found an association between higher levels of serum AFB-Ly adducts and decreased CD8⁺ cell function in human blood ¹⁶⁷. Prospective studies are needed to confirm this finding. The exact mechanisms underlying these immune toxic effects are still not clear and remain to be investigated.

Growth Impairment

Increasing evidence has shown the adverse health effects of aflatoxins on growth in humans ⁵. A large cross-sectional study demonstrated strong negative associations between levels of serum aflatoxin albumin adduct and the growth parameters (height for age, weight for age, and weight for height) in young children (9 month to 5 years) recruited in Benin and Togo, West Africa ¹⁶⁸. A longitudinal study conducted in Benin assessed the effects of aflatoxin exposure on growth of young children ¹⁶⁹. A total of two hundred children (16–37 months of age) were recruited from four villages, two with high and two with low aflatoxin exposure. The major weaning food was a maize based porridge that is frequently contaminated with aflatoxins. A strong negative correlation was found between aflatoxin-albumin adducts in serum and height increase over the 8-month follow-up period. Compared to the lowest quartile of aflatoxin-albumin adducts, the highest quartile of biomarker was associated with a mean 1.7 cm reduction in growth. Maternal exposure to aflatoxins during pregnancy was shown to affect the growth of the

first year of life in a follow-up study in Gambia ⁵⁶. The lowered birth weight was also reported to be correlated with pregnancy aflatoxin exposure in the United Arab Emirates ¹⁷⁰.

Childhood stunting is an important public health problem that contributes to approximately 1/5 of deaths among children under 5 years old in developing countries. More recently, an interest is raised on how aflatoxin exposure may affect the gut health and what is the role of aflatoxin exposure in childhood stunting. Aflatoxins that are orally ingested are mainly absorbed in the intestinal mucosa, where they may interact with intestinal epithelium and bacterial flora ¹⁷¹. However, the significance of this possible interaction remains largely unknown. In the developing world, environmental enteropathy (EE) is quite prevalent among children and may mediate stunting. EE is a subclinical condition of the small intestine and characterized by reduced intestinal absorption capacity and increased permeability. Scientists now suspect that aflatoxin exposure may play a role in the etiology of EE ¹⁷¹. However, current human evidence relevant to this issue remains scanty and further human studies as well as mechanistic studies using animal models are warranted.

Kwashiorkor

In 1935, Cicely D. Williams introduced into the scientific literature of nutritional diseases, the West African name kwashiorkor ¹⁷². Williams described that the critical period for the development of this disease is the second year of life when other foods are introduced and breastmilk is gradually replaced by a diet of foods suitable for intake and digestion by young children. The clinical appearance of hypopigmentation, hyperpigmentation and desquamation was her main focus for the differential diagnosis,

and this view is maintained to the present day. Early work by Hendrickse and others have reported the occurrence of aflatoxins in the liver or blood of children with kwashiorkor¹⁷³⁻¹⁷⁶. In several tropical countries (e.g., Nigeria, Sudan, Durban), aflatoxins have been found more frequently and in higher concentration in liver specimens from children with kwashiorkor than in controls, as reviewed by Peraica et al.¹¹⁰. These clinical observations indicate an association between aflatoxins and kwashiorkor although the causal association cannot be established. The geographical and seasonal prevalence of aflatoxins in food and of kwashiorkor shows a remarkable similarity¹⁷⁷. Clinical investigation of aflatoxin elimination in children with kwashiorkor and marasmic kwashiorkor, who were fed an aflatoxin-free diet, proved that aflatoxins in these children are slowly eliminated⁹⁸. In several studies^{173, 176, 178}, aflatoxicol was found in the liver, lung, serum, urine or stools of children with kwashiorkor, whereas this metabolite was not found in control children. It is not clear whether this difference is causally related to kwashiorkor or is a consequence of the disease.

The effects of AFB₁ on the development of kwashiorkor in mice were investigated by Kocabas¹⁷⁹. In that study, young male Swiss mice were fed either low or normal protein level diets supplemented with or without AFB₁ (0.5 mg/day). The authors found that mice given aflatoxin and a low protein diet showed decreased serum total protein and albumin levels that are a little more prominent but not statistically different from the group of mice that fed on low protein diet only, and that remarkable histopathological changes of the liver in the group fed a low protein diet and AFB₁ when compared with the group fed only a low protein diet. Their conclusion that AFB₁ could not have contributed to the development of kwashiorkor remains arguable. Krawinkel¹⁸⁰

gave an alternative explanation of this finding, "Although this finding lacks statistical significance, it indicates that aflatoxins may play a specific or unspecific role in some children. If exposure to aflatoxins is not a necessary prerequisite for the occurrence of kwashiorkor, it may contribute to the associated liver disease." Besides the weak animal evidence, there also exists discrepancy in human data. For example, Househam and Hundt ¹⁸¹ detected the urine excretion of aflatoxins in normal children from a periurban area and those hospitalized with kwashiorkor and marasmus. They found in no case was aflatoxin isolated from the urine and concluded that aflatoxin exposure is unusual in this population and that aflatoxins do not play a primary role in the pathogenesis of kwashiorkor. There are possibilities that these hospitalized children had withdrawn from aflatoxins contaminated foods and the time of urine collection could be critical for a positive finding because urinary excretion of aflatoxins usually had a short half life. Alternatively, a chance finding might be possible considering the high prevalence of aflatoxins exposure in food in these areas.

To sum up, aflatoxins can be detected more frequently and at higher mean concentrations in the sera of children with kwashiorkor than in marasmic or healthy children. Prevalence of kwashiorkor is temporally correlated with the seasonal production of aflatoxins in foods. Confirmation of the causal relationship of aflatoxins in the etiology of kwashioror will necessitate radical change in approaches to prevent and manage the disease. It has to be acknowledged that exposure to aflatoxins constitutes a greater risk to children with this disease than normal healthy children ¹⁷⁷.

Biomarkers of AFB₁

Biomarkers are cellular, biochemical, or molecular alterations that can be objectively measured in biological media, such as human tissues, cells or fluids, to indicate the exposure, effect or susceptibility in individuals¹⁸². The framework of the relationship of biomarkers to exposure and disease was brought forth by the National Research Council committee on biological markers in 1987¹⁸². This conceptual framework nicely summarizes the concept of biomarkers (Figure 2.4). Biomarkers represent a continuum of changes, and the classification of change may not always be distinct. Once exposure occurred, a continuum of biological alteration may be detected. These biological alterations serve as biomarkers of internal dose, biological effect dose, early biological effect, functional and structural changes and disease. A biomarker of susceptibility is a measurable indicator of an inherent or acquired ability of an organism to respond to the challenge of exposure. Measures of these biomarkers have great utility in the following aspects: assessing the relationship between environmental exposure and development of clinical diseases; identifying individuals at high risk for certain disease; and evaluating efficacy of intervention tools.

The use of chemical-specific biomarkers for identifying stages in the development of environmental agents related adverse outcomes can provide important information for critical public health, clinical and regulatory problems¹⁸²⁻¹⁸⁴. Since the development of a paradigm for molecular biomarkers by a committee of the National Research Council¹⁸², such chemical-specific biomarkers to environmental situations have been of great interest to scientists in different fields, in which the study of aflatoxins is a representative one. A great challenge with biomarker research is how to interpret the resulting data. The

interactions between genetic makeup and environmental factors that underlie most human diseases make the interpretation even more complex. Carefully devised strategies for validation, application, and dissemination of information about biomarkers to the public are essential for the molecular biomarker paradigm to guide public health issues. It is believed that biomarkers that indicate the mode of action of an environmental agent will be strong predictors of an individual's risk of disease. Besides, these biomarkers can more accurately classify the status of exposure of individuals, and decrease the chances of misclassification of exposure status, a problem usually encountered by epidemiological investigations. Furthermore, these biomarkers can be used to indicate the effectiveness of interventions to certain exposure or risk and clarify the interactions between chemical and other exposures such as viral infection. To sum up, the overall goals chemical-specific biomarkers development and validation are, as summarized by Groopman and Kensler¹⁸⁴:

- to assess the ambient exposures in populations and individuals;
- to improve methods in risk assessment and estimation in populations and individuals;
- to improve classification of at-risk individuals, communities and populations;
- to improve methods for assessing exposure remediation and preventive interventions;
- to further define the mechanisms of exposure-disease linkages and underlying susceptibility factors;
- and to further define the interactions of multiple agents and multiple exposure to disease outcomes.

The development of AFB₁-related biomarkers is rooted in the uncovering of the metabolism and mechanism of aflatoxin-induced hepatocarcinogenesis in animal models⁴. AFB₁ is metabolized by CYP450 enzymes to its ultimate reactive metabolite, AFB₁-8,9-epoxide. This reaction is catalyzed by CYP1A2 and 3A4 in humans⁶¹. The reactive epoxide can interact with DNA to produce an AFB₁-N⁷-guanine adduct, which can be readily detected in urine after exposure. Urinary AFB₁-N⁷-guanine adduct is an elegant biomarker of the biologically effective dose because this biomarker is promutagenic and mechanistically linked to the carcinogenicity of AFB₁⁶⁷. The epoxide is unstable and undergoes Schiff base reaction to form products that react with albumin to form albumin adducts¹⁸⁵. Aflatoxin B₁-lysine (AFB-Lys) adduct, which can be obtained after enzymatic digestion of adducted albumins, has a longer half-life and serves as a reliable biomarker of internal dose. Other oxidation metabolites, including AFM₁ and AFQ₁, are also formed. Levels of AFM₁ in urine correlate well with dietary intakes of aflatoxin in humans and serve as a biomarker of internal dose¹⁸⁶. The conjugation of glutathione with phase I metabolites produces aflatoxin B₁-mercapturic acid (AFB-NAC), a biomarker of detoxification. The more detailed information on the major biomarkers of AFB₁ is provided below.

AFB-N⁷-Guanine Adduct

AFB-N⁷-guanine is the principal DNA adduct formed in rat liver after AFB₁ exposure, which derives from covalent bond formation between C⁸ of AFB₁-8,9-epoxide and N⁷ of guanine bases in DNA. It was initially identified in *in vitro* and *in vivo* experiments in 1977²⁴ and 1978⁶⁷, respectively. The urinary excretion of this DNA adduct was reported in rat following the structural characterization. An excellent dose-

dependent correlation was found between the urinary excretion of this biomarker and AFB₁ intake over a dose range of 0.125-1.0 mg/kg⁶⁷. Approximately 1% of administered AFB₁ is bound covalently to DNA and this binding reaches a peak level at 2 h after dosing; a single dose of 25 mg AFB₁ to Fischer 344 rats resulted in a measurable level of AFB₁-DNA adducts that was more pronounced following a second daily dose¹⁸⁷. Studies on the reaction of synthetic AFB₁-8,9-epoxide with DNA *in vitro* strongly indicated adduct formation is preceded by a precovalent intercalation complex between double-stranded DNA and the highly electrophilic, unstable AFB₁-*exo*-8,9-epoxide isomer¹⁸⁸. It was also found that the *exo*-isomer of AFB₁-8,9-epoxide appears to be the only product of AFB₁ involved in reaction with DNA and reaction with N⁷ position of guanine¹⁸⁹. The availability of monoclonal antibodies specific for AFB₁-DNA adducts facilitates the measurement of this biomarker¹⁹⁰. Initial AFB₁-N⁷-guanine adduct can convert to a ring-opened formamidopyrimidine derivative, AFB₁-FAPY. The formation of AFB₁-N⁷-guanine adduct was linear over the low-dose range in all species examined and formation of which was correlated with the incidence of hepatic tumor in rat and trout¹⁹¹. Several human studies have validated the correlation between dietary exposure to AFB₁ and excretion of AFB₁-N⁷-guanine adducts in urine. For dosimetry of aflatoxin exposure, 42 individuals in the Guanxi region of China were studied¹⁹². A portion of the actual food consumed was assayed for aflatoxin content and urinary output was assayed for excretion of AFB₁-N⁷-guanine by both males and females. Through a linear regression analysis, the AFB₁-N⁷-guanine excretion and the AFB₁ intake from the previous day showed a correlation coefficient of $r = 0.65$ and P less than 0.000001. Analysis of the total AFB₁-N⁷-guanine excretion in the urine during the complete collection period plotted against

the total AFB₁ exposure in the diet for each of the individuals revealed a correlation coefficient of $r = 0.80$ and P less than 0.0000001. This urinary biomarker has been used to assess human liver cancer risk in high aflatoxin exposure areas¹³⁶, and HBV carriers have been found to have higher urinary excretion of this adduct in Taiwan¹⁹³.

AFM₁ in Urine and Milk

Holzappel et al. isolated and characterized the AFM₁ and AFM₂ in sheep urine in 1966¹⁹⁴. Before their structural characterization, several studies had reported the occurrence of an AFB₁ related compound in milk¹⁹⁵ and urine¹⁹⁶. AFM₁ was initially detected in human urine in the Philippines¹⁹⁷ and it was suggested as a potential marker to link aflatoxin exposure with human liver cancer. The good correlation between dietary AFB₁ levels and urinary AFM₁ excretion was demonstrated in a study of Guangxi Residents in China¹⁹⁸. In that study, a total of 252 urine samples were collected and analyzed by an ELISA method and the dietary intake of AFB₁ was measured on a daily basis for 1 week. The correlation coefficient was found in the range of $r = 0.82-0.86$ ($P = 0.0000$) in male population. In female population, the correlation is not significant, with a correlation coefficient in the range of 0.2-0.3 (P -values in the range of 0.26-0.53). The extrapolated values of AFM₁ excreted in the urine was in the range of 1.23-2.18% of total AFB₁ consumed for the male population and 1.30-1.78% for the female population, respectively. Another study conducted in Guangxi, China also showed a good correlation between serum aflatoxin albumin adduct levels and AFM₁ excretion in human urine ($r = 0.60$, $P < 0.00003$)¹⁸⁶.

The urinary AFM₁ biomarker has been used to indicate the risk of liver cancer. In a cohort study in Shanghai, China, it was found that people with detectable levels of

urinary AFM₁ are more likely to develop liver cancer¹³⁶. AFM₁ is of similar acute toxicity^{199,200} and about 10-fold less carcinogenic compared to AFB₁^{201,202}. On the other hand, the risk of liver cancer associated with exposure to AFM₁ (usually through milk), as estimated by Henry et al., is so low as to be immeasurable in populations with a low hepatitis B incidence (such as European populations) even in heavy consumers of milk products²⁰³. This risk assessment was carried on with two maximal contamination levels of AFM₁ at 0.05 or 0.5 µg/kg. A similar finding has been reported in Taiwan²⁰⁴. A total of 144 milk samples of three main brands in Taiwan were collected twice a month over a 1-year period. Of these milk samples, only trace amounts of AFM₁ were found (1.17-54.7 ng/L) with only one sample slightly exceeding the regulatory limit of the European Union (50 ng/L). Using a World Health Organization method of evaluating risk of liver cancer, the group at greatest risk was 6- to 9-year-old girls (average, 12.2 additional cases per billion) and the group with the lowest risk was men of 45 to 64 years of age (average, 3.45 additional cases per billion). Based on these findings, the risk for liver cancer due to ingestion of milk contaminated with AFM₁ was estimated to be low in Taiwan.

Quantitative analysis from a previous animal study has shown that approximately 1-2% of a single dose of AFB₁ is excreted as AFM₁ in the milk in cow²⁰⁵. The detection of AFM₁ in human breast milk has been reported in many different populations since the early 1980s. In certain populations, more than 80% of human breast milk samples have detectable levels of AFM₁²⁰⁶⁻²⁰⁹. Given the fact that AFM₁ is of similar acute toxicity compared to its parent compound AFB₁²⁰⁰, the presence of AFM₁ in breast milk has raised a significant health concern as infants may be more susceptible to aflatoxins than

adults. A preliminary study estimated a conversion ratio of 0.09-0.43% of AFB₁ in diet to AFM₁ in human breast milk²¹⁰.

AFB₁-Albumin Adduct and AFB₁-Lysine (AFB-Lys) Adduct

In Wistar rats that were orally exposed to a single dose of AFB₁, a linear dose-response relationship between AFB₁ dose and plasma protein binding was found in the dose range of 3.5 to 200 µg/kg body weight. Twenty-four hours after exposure, approximately 0.98-2.15% of the total dose was found to be bound to the plasma protein. In addition, a constant ratio between levels of AFB₁ binding to plasma and liver DNA was also found²¹¹. An earlier study showed that albumin is the major protein in serum that binds AFB₁ to a significant level in rat²¹², and the isolation and characterization of the major serum albumin adduct formed by AFB₁ *in vivo* in rat was demonstrated by Sabbioni et al. in 1987¹⁸⁵. That study supported a structure in which the terminal dihydrofuran ring of AFB₁ has been converted to a pyrrolinone ring, and proposed that the initial adduct is formed by condensation of the dialdehyde tautomer of 8,9-dihydro-8,9-dihydroxy-AFB₁, with the ε-amino group of lysine, to form a Schiff base, and that the Schiff base undergoes an Amadori rearrangement to an α-amino ketone (Figure 2.5). An alternative structure of the lysine adduct was also proposed²¹³, which requires an alternate scheme because the formation of an amide is not particularly feasible following an Amadori rearrangement (Figure 2.6). The relative contribution of AFB₁-8,9-epoxide and AFB₁ dialdehyde in formation of lysine adduct has been discussed (Figure 2.7)²¹⁴. It is noted that the apparent rate of reaction of AFB₁ dialdehyde with *N*²-acetylLys is approximately 7 times greater than for the epoxide at pH 7.2. In addition, the rapid, irreversible reaction of the epoxide with H₂O ($k = 42 \text{ min}^{-1}$) suggests that most of the Lys

adducts formed might be from the reaction with the dialdehyde. Using the kinetic model (Figure 2.7), it is estimated that ~ 90% of the lysine adduct would arise from the dialdehyde and ~ 10% from the epoxide in the absence of other reactions at a concentration of 1 mM (*N*²-acetyl) lysine.

The radioactivity associated with serum albumin following administration of [¹⁴C]AFB₁ to rats was cleared with a half-life of 2.5 days¹⁸⁵. Similarly, the half life of serum AFB-Lys adduct has been reported to be 55 h²¹¹ and 2.6 days²¹⁵ in rat. The elimination of serum AFB-Lys followed first-order kinetics. Compared to the short half life of urinary aflatoxin biomarkers, the serum AFB₁ albumin adduct in humans has a biological half life of 21 days, making it a good biomarker for integrating human exposure for several weeks or even months¹⁹². Because the plasma protein binding requires metabolic activation, this adduct could be used as a biomarker of effective dose.

A good correlation of AFB-Lys adduct with AFB₁-DNA adduct has been reported in several animal species^{211, 216}, including rat, Guinea pig, hamster, and mouse. Wild et al. assessed the aflatoxin B₁ albumin adduct as a basis for comparative carcinogenesis between animals and humans²¹⁶. Three strains of rat (Fischer 344, Wistar, and Sprague-Dawley), and one strain each of guinea pig (Hartley), hamster (Syrian golden), and mouse (C57BL) were treated by gavage with up to 14 daily doses of between 1 and 80 μg AFB₁/kg body weight. A dose response in both AFB₁-albumin and AFB₁-DNA adducts was seen for all species and strains with steady-state adduct levels at 14 days. The level of both adducts was in the following order: rat > guinea pig > hamster > mouse. The levels of AFB₁-albumin adduct also reflect at least qualitatively the relative susceptibility of the different species to AFB₁ carcinogenesis; the rat is sensitive and the hamster and

mouse are resistant. In addition, when the levels of the albumin and DNA adducts at 14 days were plotted against each other for all species and strains, a correlation was observed ($r = 0.83$, $P \leq 0.0001$; $n = 57$; two-tailed test), suggesting a constant relationship between the level of binding of AFB₁ to serum albumin and liver DNA. The level of AFB₁-albumin adduct formed as a function of a single dose of AFB₁ in rodents was compared to data from humans exposed environmentally to AFB₁. The yield (pg AFB-Lys equivalent/mg albumin/ μg AFB₁/kg body weight) for the three rat strains after a single dose was between 0.3 and 0.51, and a value for the mouse of < 0.025 . At human exposure level, a linear relationship between formation of AFB-Lys adduct and liver AFB-DNA adduct has been demonstrated in rats (0.16 ng/kg - 12.3 $\mu\text{g}/\text{kg}$)²¹⁷. These data suggest that the level of serum AFB-Lys adducts is a useful biomarker for AFB₁ dosimetry and may reflect the DNA adduct levels in the target tissue. In a human study in Guangxi, China, serum AFB₁ albumin adducts were correlated with dietary AFB₁ intake and urinary excretion of AFM₁. Regression analysis revealed that 1.4-2.3% of ingested AFB₁ is covalently bound to serum albumin¹⁸⁶, which is similar to that found in rats. AFB-Lys adduct, under proper storage conditions, is stable in sera for more than 20 years²¹⁸. In those populations at high risk for aflatoxin exposure such as southeast Asia and sub-Saharan Africa, usually more than 90% of the human populations have detectable AFB-Lys adducts in their sera. Our data from Guangxi, Burkina Faso, Ghana, and Uganda also support this finding.

This biomarker has been widely applied in epidemiological studies to assess the exposure to and risk of aflatoxin exposure in high-risk populations^{134, 219} and to test the efficacy of different intervention approaches^{220, 221}. During the 2004 outbreak of

aflatoxicosis in Kenya, this biomarker was used to establish the exposure to aflatoxins in affected individuals¹¹⁸. Currently, the most commonly used methods to measure this adduct are the ELISA²²² method, the immunoaffinity column or solid phase extraction (SPE) based HPLC method with fluorescence detection²²³, although other methods are still available, such as isotope dilution mass spectrometry (IDMS) method^{215, 224}, and radioimmunoassay (RIA)^{186, 219}. The results of antibody-based assays are influenced by the specificity and the sensitivity of the antibodies used. Since the major AFB-albumin adduct was identified as the AFB-Lys adduct^{185, 213}, the development of more specific monoclonal antibodies recognizing this adduct was initiated and became available in 2001²²⁵. The detection limit of HPLC and IDMS method is comparable, that is, approximately 0.5 pg/mg albumin. The ELISA method usually gives higher value as compared to the IDMS method (2.6-fold difference) but the values are well correlated ($r = 0.88$, $P < 0.0001$)²²⁶. The result suggests that ELISA method may result in nonspecific binding of the antibody to other aflatoxin adducts other than AFB-Lys adduct. For this reason, the data generated from the same method are favored for comparative purposes.

AFB₁-Mercapturic Acid (AFB-NAC)

Biological effects of AFB₁ are principally induced by the metabolite, AFB₁-8,9-epoxide. Detoxification of the epoxide can be mediated in part by glutathione S transferase (GST)⁷². The enzymatic conjugation of the AFB₁-epoxide with glutathione is a mechanism of protection against its toxic effects. Under physiological conditions, the glutathione conjugates undergo further metabolic processing to form mercapturic acids, which are expected to be excreted in urine and have been tested as a biomarker of AFB₁. Scholl et al.²²⁷ synthesized and characterized aflatoxin B₁-mercapturic acid (AFB-NAC)

and investigated the dose-response relationship between AFB₁ exposure and urinary excretion of this conjugate in rat. They found that approximately 1% of AFB₁ dose is excreted as AFB-NAC within 24 h exposure in F344 rats. This conjugate is excreted in urine as a linear function of dose over the range of doses 0.03-0.8 mg/kg body weight. The finding that AFB-NAC is dose-dependently excreted in urine provides the basis for investigating its applicability as a biomarker of GST induction in aflatoxin chemoprevention studies²²⁸.

In a randomized, placebo-controlled, double-blind phase IIa chemoprevention trial, the chemopreventive effects of oltipraz was tested in residents of Qidong, People's Republic of China²²⁹. The results showed that a daily intervention with 125 mg oltiprza lead to a 2.6-fold increase in median AFB-NAC excretion compared to the placebo group. This finding highlights the feasibility of inducing phase II enzyme as a chemopreventive strategy in humans. In another study evaluating the efficacy of green tea polyphenols (GTPs) in modulating AFB₁ biomarkers, a total of 352 urine samples were collected from a 3-month chemoprevention trial with 500 mg GTPs, 1000 mg GTPs and a placebo treatment²²⁰. Significant elevations in median AFB-NAC levels were found in both 500 and 1000 mg groups compared with the placebo group at both 1 month ($P < 0.001$) and 3 months ($P < 0.001$) of GTPs intervention. This finding suggests that GTPs effectively modulate AFB₁ metabolism and enhances the metabolic detoxification of AFB₁ and holds promise as a potential intervention tool in human studies.

Mutation at Codon 249 of Tumor Suppressor Gene *p53*

The mutations in the *p53* tumor suppressor gene is more frequently detected in HCC cases residing in high AFB₁ exposure regions than those in low AFB₁ exposure

regions (50% vs 20%), and the spectrum of mutations is quite different²³⁰⁻²³³. Among those HCCs from high AFB₁ region, more than half of the tumors contain an AGG to AGT missense mutation in codon 249 (AGG to AGT) of the *p53* gene, leading to the replacement of arginine by serine. High prevalence of mutations at codon 249 of the *p53* gene in HCCs was also found in Senegal²³⁴, this typical mutation was detected in 10 of the 15 tumor tissues tested (67%). Although the number of cases is not large, this frequency of mutation in codon 249 of the *p53* gene is the highest described. A study of Mexico HCC cases showed that 7 of 16 HCC cases (44%) were found to have AGG to AGT mutations at the exon 7 of *p53* gene codon 249²³⁵, in which the serum AFB₁-albumin adduct levels were found in a range of 0.54 to 4.64 pmol/mg albumin.

It has been demonstrated that AFB₁ induces the G to T transversions in the third position of codon 249 of the *p53* tumor suppressor gene in human HCC cells HepG2²³⁶. Different cellular *in vitro* models that take into account species and tissue specificity have been developed to examine the contribution of different metabolic enzymes in inducing AFB₁ associated *p53* gene mutation. A non-tumorigenic SV40-immortalized human liver epithelial cell line (THLE cells) that retains most of the phase II enzymes, but has markedly reduced phase I activities was used for stable expression of the human CYP1A2, CYP2A6, CYP2B6 and CYP3A4 cDNA. The four genetically engineered cell lines (T5-1A2, T5-2A6, T5-2B6 and T5-3A4) were developed, which produced high levels of the specific CYP450 proteins with comparable or higher catalytic activities compared to that in human hepatocytes. At similar levels of total DNA adducts, both the T5-1A2 and T5-3A4 cells showed AGG to AGT transversions at codon 249 of the *p53* gene²³⁷.

The mechanisms underlying AFB₁-induced G to T transversion in codon 249 of the *p53* gene mutation remains largely unknown. The cytosine residues of CpG sites in the coding region of the *p53* gene are known to be methylated in a variety of human tissues²³⁸. It has been hypothesized that mutational hot spots at methylated CpG sequences may represent preferential carcinogen targets²³⁹. An *in vitro* study was designed to test the role of cytosine methylation on the mutation spectrum using AFB₁ as the carcinogen. A wild type *p53* cDNA fragment was cloned into a pUC18 vector to generate a DNA template for AFB₁ treatment and the mutations were identified by DNA sequencing. The results showed that cytosine methylation enhances AFB₁-induced guanine mutations at CpG sites but no mutations were detected at codon 249²⁴⁰. This finding suggests that the DNA sequence at the third base of codon 249 itself may be insufficient to be a hot spot target.

Biomarkers of exposure to AFB₁ in human populations

The currently favored method of measuring human exposure consists of the analysis of body fluids for the presence of aflatoxin biomarkers^{241, 242}. Each biochemical process results in derivatives that have a characteristic half-life within the body, and thus the measurement of these derivatives can reflect the exposure over a period of days, weeks, or months. Recent exposure to aflatoxin is reflected in urine as directly excreted AFM₁ and other detoxification products, but only a small fraction of the dose is excreted in that way. Measurements of aflatoxin and its byproducts in urine have been found to be highly variable from day to day, which reflects the wide variability in the contamination of food samples, and, for this reason, the measurement of AFM₁ on a single day may not be a reliable indicator of a person's chronic exposure²⁴¹⁻²⁴³. The aflatoxin-albumin

adduct that can be measured in peripheral blood and has a half-life 21 days in humans¹⁸⁵. This biomarker could integrate the exposure over a longer period and hence is a more reliable indicator of a person's chronic exposure. However, it should be remembered that the fraction of the ingested aflatoxin processed into any particular metabolite is variable. A given concentration of any particular biomarker cannot be used to make assumptions about the total dose or the amounts directed into any other competing pathway, partially due to the individual phenotype in metabolic enzymes as discussed in biotransformation section. Human exposure data using biological markers as a measure are relatively rare, because the locations of such measurements are mainly in West Africa^{128, 130, 168, 178, 244-246} and China²¹⁹. However, there are increasing data showing a high detection rate of aflatoxin biomarkers in Malaysia²⁴⁷, Thailand²⁴⁸, Egypt²⁴⁹, Haiti, Uganda, Burkina Faso (unpublished data), and the United States²⁵⁰ (Table 2.3). These data also demonstrate major variations in seasonal exposure^{128, 245}, with summer months usually have higher exposure than winter months²⁰⁸, which reflects the natural development of contamination in storage. These biomarker data show that, regardless of food preparation practices, the human populations of these developing countries are widely and significantly exposed to aflatoxin. Given that little is done to decontaminate foods in many developing countries, it is likely that the prevalence of chronic exposure in these countries is similar to that measured in these studies with the use of biomarkers².

Intervention strategies for reducing aflatoxins exposure and risk

It is estimated that approximately 4.5 billion people in the developing world are at risk for chronic exposure to aflatoxins². Intervention strategies for reducing aflatoxins exposure or risk have been the concerns of several reviews^{2, 4, 27, 28}. These strategies

could be undertaken at different stages, including pre-harvest (reduce toxin production and bio-control), during food processing and storage (improve environmental setting, selection, and education), and after ingestion (reduce active metabolite formation and bioavailability). Vaccination of infants against HBV and HCV to remove the interaction with AFB₁ exposure is of priority to reduce human HCC risks in the long-term¹⁴¹. The evaluation of different intervention strategies has been defined as one of the four gaps (quantification of health effects and disease burden of aflatoxin exposure; evaluation of current intervention strategies and disseminate the result; disease surveillance, food monitoring, laboratory and public health responses; development of response protocols addressing aflatoxicosis) for future effort by the organized workshop that was held after the outbreaks of aflatoxicosis in Kenya²⁷.

Primary intervention strategies for reducing aflatoxins exposure include post-harvest intervention, using trapping agents to reduce absorption such as clay and chlorophyllin. A strategy of post-harvest intervention has been reported in a community-based study in Kindia region of Guinea, West Africa²⁵¹. A 60% reduction of mean aflatoxin level was achieved after 5-month in the intervention group, compared to the control. Reduction of the percentage of persons with detectable level of aflatoxin albumin adducts were also found. In Haiti, results have been achieved to reduce aflatoxin contamination in market peanuts, through visual-tactile sorting of the shell, flotation of select kernels, immediate roasting and grinding, and monitoring²⁵². The level of AFB₁ in peanuts showed a reduction from 412.5 ± 32.1 ppb in September 2006 to 125 ± 7.1 ppb in November 2006 and further to a level of 0.20 ± 0.10 ppb in January 2007. Therefore, primary intervention is a promising approach to reduce aflatoxin exposure, and several

primary intervention strategies in West Africa have been determined to be cost effective²⁵³.

Chemoprevention is a strategy that involves the use of natural or synthetic agents to block, retard, reverse or modulate the carcinogenic process of environmental agents²⁵⁴. Aflatoxin biomarkers were first used as surrogate endpoints in a phase II chemoprevention trial in Qidong, China^{229, 255, 256}. Individuals receiving 500 mg oltipraz once a week for 8 weeks had a significant longitudinal decline in aflatoxin-albumin biomarker levels beginning 1 month into the intervention and continuing for 1 month after treatment was stopped. An association was observed within participants in the 500 mg group between rate of decline of aflatoxin-albumin adducts and excretion levels of AFM₁. AFB₁-NAC levels were elevated 2.6-fold in the 125 mg group, compared to placebo control. Oltipraz seems to regulate both phase I and phase II metabolizing enzymes to modulate the aflatoxin metabolic pathways²⁵⁷. The transcription factor Nrf2 mediated signal pathway appears to be essential for the induction of phase 2 enzymes by oltipraz. While oltipraz significantly reduced multiplicity of gastric neoplasia in wild-type mice by 55%, there is no effect on tumor burden in *nrf2*-deficient mice²⁵⁸. Chlorophyllin, a water-soluble derivative of chlorophyll, has demonstrated its anticarcinogenic properties in a number of animal models^{259, 260}. The primary mode of action of chlorophyllin appears to be sequestration of aflatoxin through complexation in a 1:1 ratio²⁶¹, although experimental data have characterized its enzyme-inducing properties that may also contribute to its mechanism of action²⁶². The effect of chlorophyllin on disposition of aflatoxin has been assessed in a randomized, double-blind, placebo-controlled chemoprevention trial in residents of Qidong, China. One

hundred and eighty healthy adults were randomly assigned to ingest 100 mg of chlorophyllin or a placebo three times a day prior to each meal for 4 months. Urine samples were collected from both groups to test for the modulation of urinary aflatoxin N⁷-guanine, a primary endpoint. Aflatoxin-N⁷-guanine could be detected in 105 of 169 available samples. Chlorophyllin consumption at each meal led to an overall 55% reduction ($P = 0.036$) in median urinary levels of this aflatoxin biomarker compared with those subjects taking placebo. This study provides evidence that dietary supplementation with chlorophylls or ingestion of foods rich in chlorophylls may represent practical means to prevent the development of HCC in people chronically exposed to aflatoxins.

The natural product NovaSil (NS) clay has been shown to be safe in animals and human subjects ^{263, 264}. When included in the diet, NS clay has been shown to prevent aflatoxicosis in a variety of animals and epidemiological studies have demonstrated the efficacy of NS clay in reducing the serum biomarkers of AFB₁ in populations exposed to aflatoxins through contaminated foods ^{29, 221}. Green tea polyphenols (GTPs) have demonstrated efficacy in cancer prevention in rat models and reduction of oxidative DNA damage biomarkers in clinical trials ^{265, 266}. The induction of phase II enzyme and scavenging of oxidative damage might account for the protective effects in aflatoxin exposed population ^{220, 266}. When reviewing the current research on NS clay and aflatoxins, Phillips et al. concluded that the aflatoxin sequestering clays should be rigorously evaluated *in vitro* and *in vivo* and several criteria should be met: favorable thermodynamic characteristics of mycotoxin sorption; tolerable levels of priority metals, dioxins/furans and other hazardous contaminants; safety and efficacy in multiple animal species; safety and efficacy in long-term studies; and negligible interactions with

vitamins, iron and zinc and other micronutrients ²⁹. Most of the recommended criteria could be conveniently applied to the research of other chemopreventive agents as well. The readily accessible aflatoxin biomarkers and analytical tools greatly facilitate the evaluation of different intervention strategies in both animal experiments and human epidemiological studies.

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Table 2.1. Acute toxicity of AFB₁.

Species	Strain	LD ₅₀ (mg/kg)	Sex	Age or weight	Route
Duck	Khaki-Campbell	0.36	M,F	1 day	P.O.
Duckling	Peking	0.34	M,F	1 day	P.O.
Chicken		18	M	21 days	P.O.
Turkey	Beltsville	3.2	M	15 days	P.O.
Trout	Mt. Shasta	0.81	M,F	9 months	I.P.
Catfish	Channel	11.5	M,F	9.3-0.5 kg	P.O.
Mouse	Swiss	1.5	M,F	newborn	P.O.
	Swiss	>150	M	30 days	I.P.
	Swiss	40	M	58 days	I.P.
	Swiss	12	M	100 days	I.P.
	CD-1	7.3	M	weaning	P.O.
Rat	Porton	6.25	M	42 days	P.O.
	Porton	18	F	43 days	P.O.
	Fischer	4.2	M	42 days	I.P.
	Fischer	1.1	M,F	2 days	S.C.
	Fischer	8	M,F	21 days	I.P.
Hamster	Syrian	12.8	M	30 days	P.O.
	Syrian	5.85	F	42 days	I.P.
Guinea Pig		1	M	56 days	P.O.
		1.8	F	56 days	P.O.
Rabbit	Dutch breed	0.3	M,F	90 days	I.P.
Cat	Mixed breed	0.55	M,F	adult	P.O.
Dog	Mixed breed	0.8	M,F	weaning	P.O.
Pig	Poland China	0.62	M	weaning	P.O.
Sheep	Cross breed	2	M	2 years	P.O.
Baboon	Wild	2.2	M	adult	P.O.
Monkey	Cynomolgus	2.2	M	adult	P.O.
	Macaque	8	F	adult	P.O.

I.P., intraperitoneal; P.O., per oral. S.C., subcutaneous (References #99, 101-104)

Table 2.2. Acute aflatoxicoses in humans.

Time	Country	# of Subjects affected	Contaminated foods	Reference
1966	USA	1	purified mixtures containing	117
1970	Uganda	1	Cassava	116
1974	Western India	397	maize	111
1974	Northwest India	994	maize	112
1982	Kenya	20	maize	113
1988	Malaysia	13	a Chinese noodle, "loh see fun"	114
2004	Kenya	317	maize and maize products	118

Table 2.3. A compilation of data on biomarkers of exposure to aflatoxin in humans

Country	Exposure rate (%)	Range or test type	Reference
Benin	99	5–1064 pmol/mg albumin	Gong et al (168)
China (Guangxi province)	89	0.9–3569 pg/24-h urine	Wang et al (219)
Egypt	65.3	AFM ₁ in breast milk, 0.2 to 19.0 ng/mL	Tomerak et al (249)
Gambia	95	0–720 pmol/mg albumin	Allen et al (128)
Ghana	90.7	0.44 - 286.73 pg/mg albumin	Shuaib et al (244)
Guinea	90	0–385 pmol/mg albumin	Diallo et al (130)
Malaysia	98	0.81 - 13.67 pg/mg albumin	Leong et al (247)
Nigeria	40-90	Lung autopsies	Oyelami et al (178)
Sierra Leone	95–99	Aflatoxin in urine	Jonsyn-Ellis (245)
Sudan, Zimbabwe, Ghana, Liberia, Kenya, Transkei	32	Aflatoxin in urine	Hendrickse et al (246)
	99	Liver biopsy of kwashiorkors	
Thailand	44	AFM ₁ in breast milk	Hollstein et al (248)
United States (San Antonio and Lubbock)	15	0.52-16.01 pg/mg albumin	Johnson et al (250) and unpublished data
Burkina Faso	100	0.56 - 918.72 pg/mg albumin	unpublished data
Haiti	67.5	0.41 - 130.39 pg/mg albumin	unpublished data
Uganda	95.3	0.41 - 253.11 pg/mg albumin	unpublished data

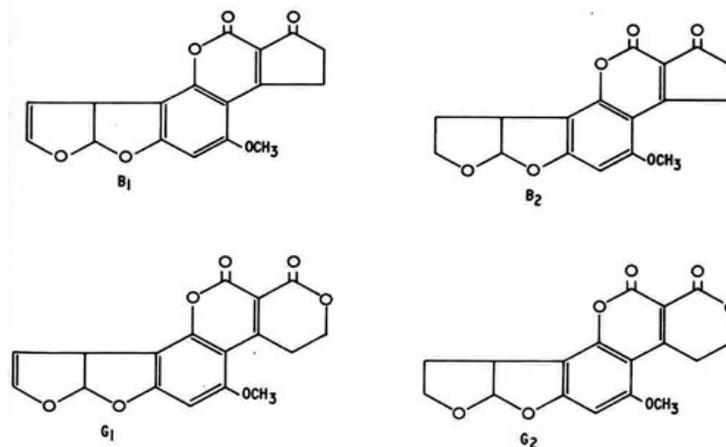


Figure 2.1. Structures of aflatoxin B₁, B₂, G₁, and G₂

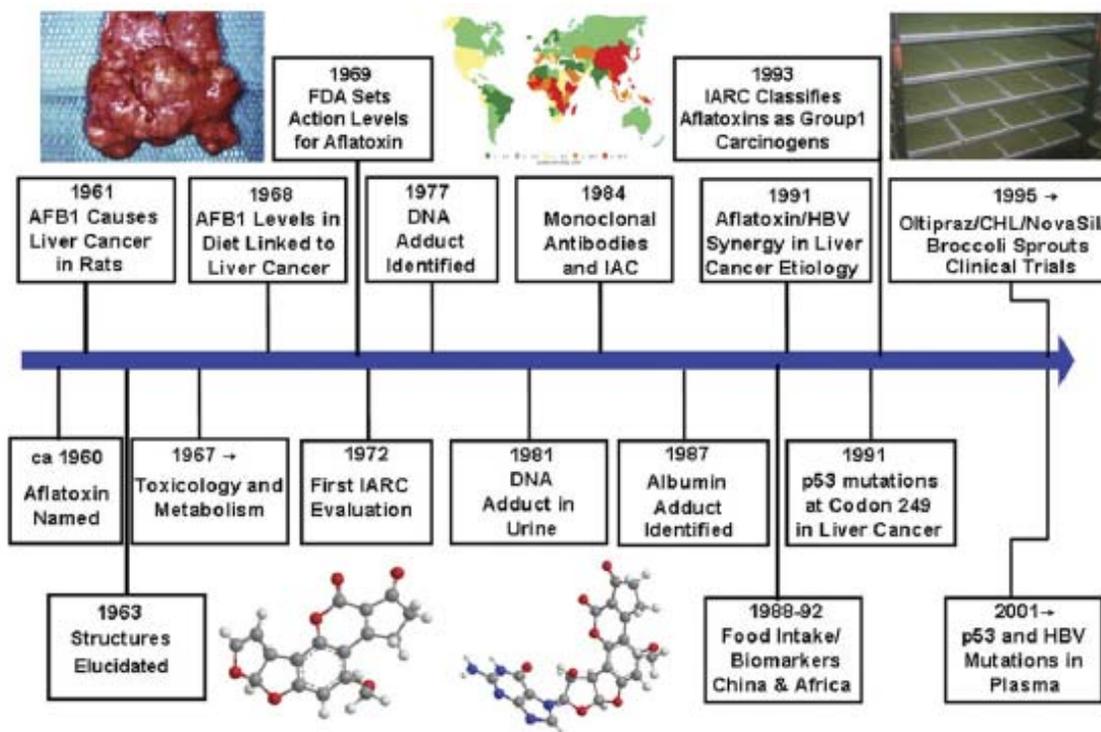


Figure 2.2. Time line for key events in aflatoxin research (Reference #4)

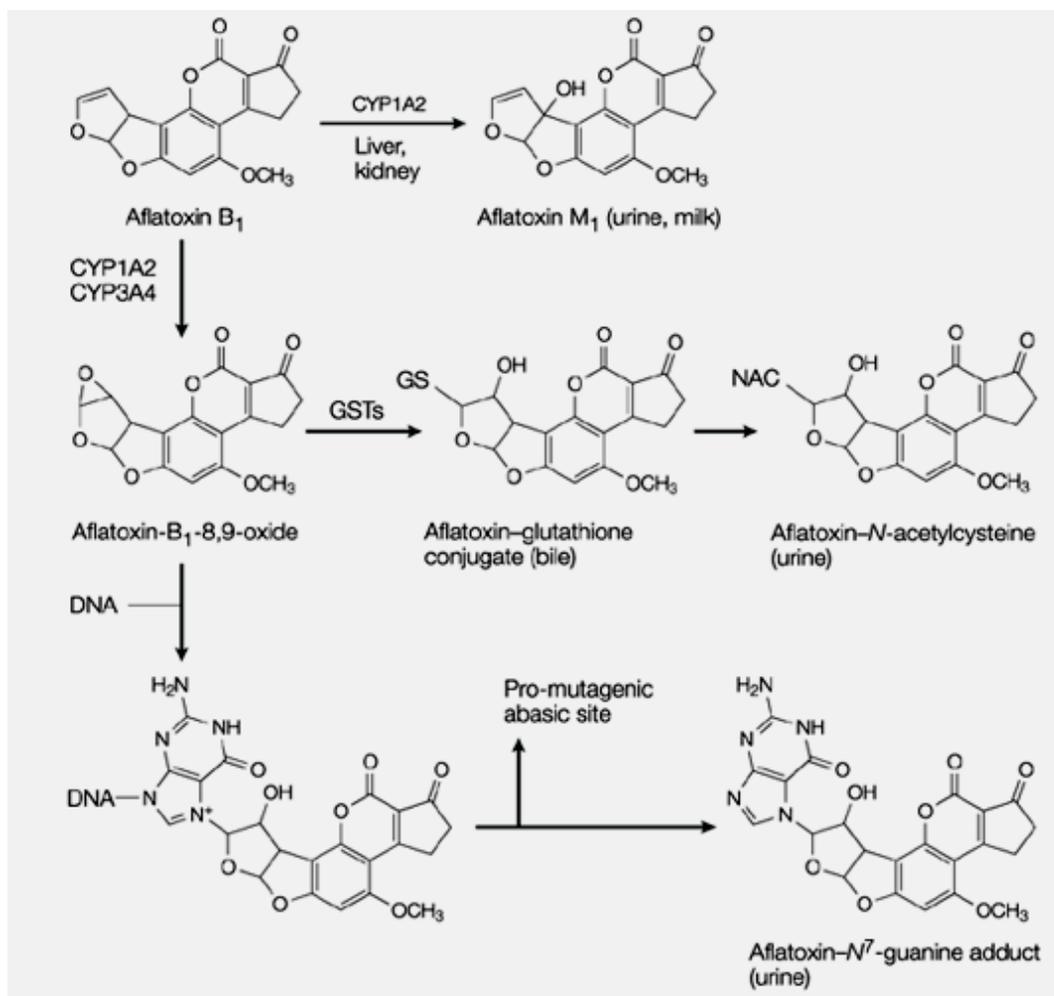


Figure 2.3. Detoxification pathways of aflatoxin B₁ (Reference #73)

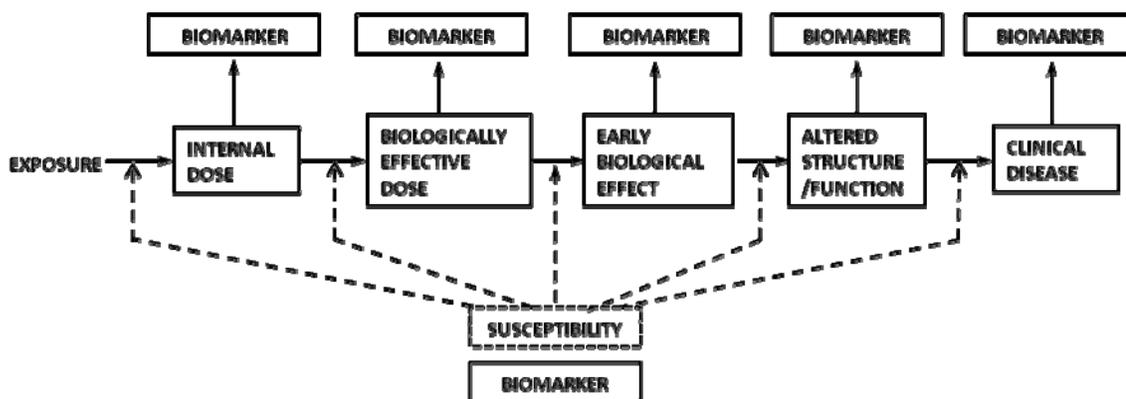


Figure 2.4. Biomarker paradigm (Reference #182)

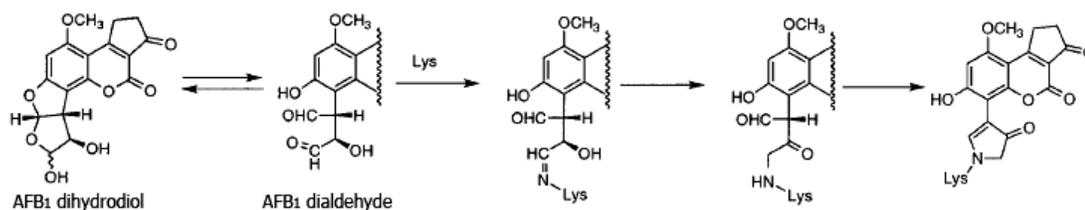


Figure 2.5. Proposed reaction of AFB₁ dialdehyde with lysine (Reference #213)

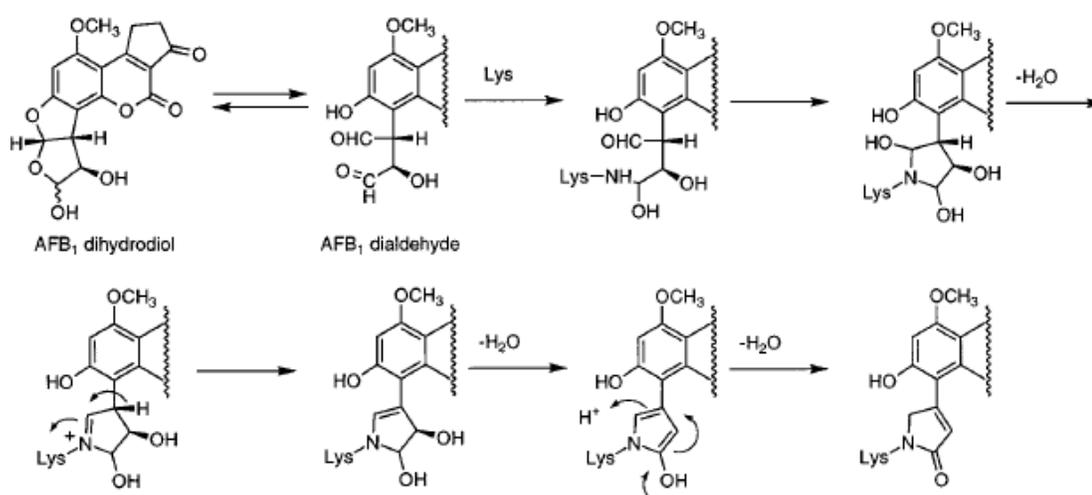


Figure 2.6. Alternative reaction of AFB₁ dialdehyde with lysine (Reference #214)

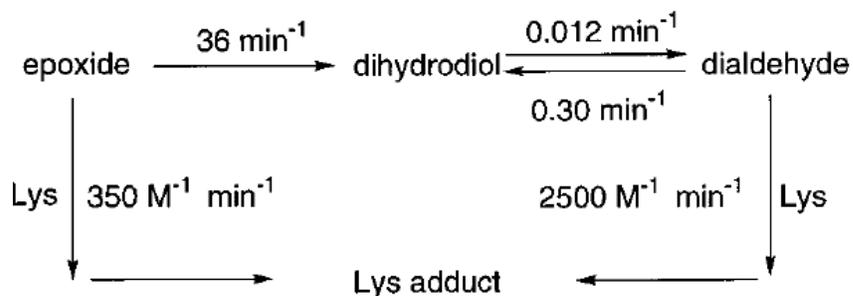


Figure 2.7. Proposed kinetic model for relative contribution of AFB₁-8,9-epoxide and AFB₁ dialdehyde in formation of lysine adduct (Reference #214)

CHAPTER 3

DEVELOPMENT OF A HPLC WITH FLUORESCENCE DETECTION METHOD

FOR MEASURING SERUM AFLATOXIN B₁-LYSINE ADDUCT ¹

¹ Qian, G. Tang, L. Liu, W. and Wang, J.-S. To be submitted to *Toxicology*.

Abstract

Aflatoxins (AF) exposure is a major public health problem in the developing world, because acute aflatoxicosis occurs frequently and chronic exposure is a major risk factor for human liver cancer. Serum aflatoxin B₁-lysine adduct (AFB-Lys) is a reliable biomarker of AF exposure. The aim of this study is to develop a solid-phase extraction based high pressure liquid chromatography (HPLC) method with fluorescence detection for rapid detection of serum AFB-Lys. The protocols were extensively optimized, including conditions of enzyme digestion; comparison of various commercial solid-phase extraction cartridges, and for HPLC separation and quantification. The optimal level of AFB-Lys was detected in serum samples digested with pronase at 37°C for 3 h. Among various cartridges, OASIS MAX™ gave better recovery and efficiency, as compared to OASIS HLB™ and Sep-Pak™ cartridges. Calibration curves were linear for concentrations ranging from 0.01-5.0 ng/mL. The method was tested with AFB₁ treated rat serum and normal human serum spiked with synthesized authentic AFB-Lys from low to high concentrations. The imprecision based on 15 measurements on 3 separate days was 5.93% and 2.87% for low quality control (LQC) and high quality control (HQC) samples, respectively. The inaccuracy for repeated measurements was 4.97% and 4.44%, for LQC and HQC, respectively. Recoveries ranged 72% to 95% based on various adduct concentrations. Under the current procedures, AFB-Lys was stable for up to 12 h and decreased about 10% at 24 h at room temperature. Serum AFB-Lys levels had no

significant changes for up to 3 freeze-thaw cycles. The limit of detection was 0.4 pg/mg albumin.

Key words: aflatoxin; aflatoxin B₁ lysine; biomarker; HPLC.

Introduction

Aflatoxins are a large group of secondary fungal metabolites produced primarily by *Aspergillus flavus* and *Aspergillus parasiticus*, which occur naturally in foods such as corn, peanuts and oilseeds ¹. The human diseases caused by aflatoxins remain a largely ignored public health issue ². Human epidemiological studies have recognized aflatoxin exposure as a major risk factor for human hepatocellular carcinoma (HCC) in southeastern Asia and Sub-Saharan Africa ³. It is estimated that approximately 4.5 billion people are affected by chronic exposure to aflatoxins in the developing world ⁴.

The development, validation and application of molecular chemical-specific biomarkers have the advantage of assessing internal human exposure to environmental agents on an individual basis, and therefore more accurate and precise than traditional food sampling survey ⁵⁻⁷. Increasing knowledge of the toxicology and mechanisms of aflatoxins toxicity have provided insights into measurable biomarkers that have gained practical application in many human populations for assessment of internal human exposure and efficacy of intervention strategies ⁷⁻⁹.

Aflatoxin B₁ (AFB₁) is the most potent among naturally occurring aflatoxins, its mutagenicity and carcinogenicity, have been proven by numerous *in vitro* and *in vivo* test models and human studies as well ¹⁰. AFB₁ is metabolized and activated by several cytochrome P450 (1A2, 2A6, 3A4, 3A5) to its most potent metabolite, *exo*-8,9-epoxide ¹¹⁻¹³. This active metabolite can bind to protein and DNA, and the ratio of hepatic protein to DNA adducts is constant even at very low dose ¹⁴. The epoxide can readily attack

DNA to form the AFB₁-N⁷-guanine adduct¹⁵ if it has not been hydrolyzed or conjugated with glutathione¹⁶. The AFB₁-N⁷-guanine adduct is excreted in urine and represents a short term effect marker with a half-life of about 7.5 hours in human¹⁷⁻¹⁹. Albumin is the only protein in serum that binds AFB₁ to a significant level in rats and about 1-3% of a single dose of AFB₁ is bound to albumin in rats^{14, 20, 21}. AFB₁ albumin adduct represents a relative long-term exposure marker which has a half-life of 2.5 days in rats and 21 days in humans^{22, 23}. The isolation and characterization of serum AFB-Lys adduct *in vivo* was reported by Sabbioni et al.²², and further discussed by Guengerich et al. who examined the kinetics of reaction of AFB₁ dialdehyde and AFB₁-8,9-epoxide with lysine *in vitro*²⁴. Serum AFB-Lys has been employed as an exposure biomarker of AFB₁ in many human epidemiological studies²⁵⁻³¹.

Since the late 1980s, different analytical methods have been developed for quantification of AFB-Lys in human sera, including radioimmunoassay (RIA)^{32, 33}, enzyme linked immunosorbent assay (ELISA)³⁴, immunoaffinity column based high performance liquid chromatography (HPLC) method with fluorescence detection^{27, 35}. The isotope-dilution mass spectrometry (IDMS) method has also been developed for measuring AFB-Lys adduct in human and rat sera^{21, 36, 37}. However, the use of antibodies in sample purification or detection determines that the sensitivity and specificity of these methods are highly dependent on the antibodies selected. The prerequisite of LC/MS equipment and use of isotope in IDMS method makes it not particularly feasible in developing countries which lack such resources. In this study, we presented a non-

antibody based method for measuring serum AFB-Lys adducts using solid-phase extraction followed by HPLC separation with fluorescence detection.

Experimental Procedures

Chemicals and standards

Unlabeled AFB-Lys was synthesized using previously reported methods^{22, 38}. Normal human serum, protein standard, ammonium phosphate monobasic, formic acid (FA) and phosphate-buffered saline (PBS) were obtained from Sigma Aldrich Chemical Co. (St. Louis, MO). Pronase (25 kU, nuclease-free) was purchased from Calbiochem (San Diego, CA). Albumin standard was obtained from Pointe Scientific Inc (Canton, MI). Methanol (MeOH), acetonitrile (ACN) and water were purchased from Mallinckrodt Baker Inc. (Phillipsburg NJ) as HPLC grade. Ammonium hydroxide was purchased from EMD Chemicals (Gibbstown, NJ). Protein assay dye reagent concentrate and protein standards were purchased from Bio-Rad Laboratories Inc. (Hercules, CA). Albumin reagent was purchased from Thermo Scientific Inc. (Waltham, MA). All other reagents or solvents are of highest grade available.

Equipment and Other materials

The Agilent 1100 series HPLC system (Santa Clara, CA) was used to analyze AFB-Lys and DU800 Series UV/Vis Spectrophotometer from Berkman Coulter Inc. (Fullerton, CA) was used to measure serum total protein and albumin content. Centriva Concentrator and Centriva coldtrap from Labconco Corporation (Kansas city, MO) were used for vacuum drying the purified eluates. VX-2500 multi-tube vortexer from VWR

International (West Chester, PA) were used to mix reconstituted samples. Oasis MAX™ 1cc (30 mg) extraction cartridge, Oasis HLB™ 3cc (60 mg) cartridge and Sep-Pak C18™ cartridge were purchased from Waters (Milford, MA). ZORBAX Eclipse XDB-C18 column (4.6×250 mm, 5 micron) was bought from Agilent Technologies Inc (Santa Clara, CA).

Quantification of Synthesized Standard

The synthesized AFB-Lys was quantified by measuring the absorbance at 400 nm wavelength in MeOH and the published molar extinction coefficient of 30866 ($\text{Lmol}^{-1}\text{cm}^{-1}$), as described in a previous study³⁸.

LC/MS/MS Confirmation of AFB-Lys Standard

The confirmation of synthetic AFB-Lys standard was done using a ThermoFinnigan HPLC system with surveyor autosampler and pump (Thermo Electron Corp. San Jose, CA, USA). The analytical column was a Thermo Betabasic-18 (150×2.1 mm, 5 μm). The mobile phases were: A, 10% ACN with 0.3% formic acid (FA); B, 90% ACN with 0.3% FA. The autosampler compartment was maintained at 4°C and column at 35°C. The mobile phase was run as a gradient at a flow rate of 0.2 mL/min. Initially, the mobile phase consisted of 100% A. Then 100% A was linearly decreased to 0% in 15 min. Isocratic elution was maintained for 10 min before changing back to the initial condition in 1 min and was the column was re-equilibrated to the initial condition for another 9 min. The single run-time was 35 min. The eluate was introduced into a ThermoFinnigan LCQ Advantage tandem mass spectrometer with electrospray

ionization. The positive ion mode was employed for selective reaction monitoring (SRM) m/z transition of 457 ($M+H^+$) to 394 for AFB-Lys³⁹: capillary temperature 300°C, source voltage 4 kV, capillary voltage 7 V, the collision energy was set at 40 eV, sheath gas flow 60 units, auxiliary gas 20 units.

Binding Efficiency of Commercially Available Cartridges

MAX, HLB and Sep-Pak cartridges were compared for their binding efficiency, as shown in Figure 3.1. Briefly, the MAX cartridge was pre-conditioned with 1 mL of MeOH and equilibrated with 1 mL of water. Then 150 μ L serum diluted with 500 μ L DI water was loaded into the cartridge, followed by sequential washing with 1 mL water twice, 1 mL 70% MeOH, 1 mL 1% ammonium hydroxide in MeOH and 0.5 mL MeOH at a speed of approximately 1 mL/min. Vacuum was used to dry the cartridge after the last 0.5 mL MeOH washing. Elution of the AFB-Lys was done with 1 mL 2% formic acid in MeOH into a labeled 12 \times 75 mm glass tube. Same procedure was used on both HLBTM and Sep-PakTM cartridges. Briefly, the cartridges were conditioned with 1 mL MeOH and equilibrated with 1 mL water. Serum (150 μ L) was diluted with 500 μ L water which was passed through the cartridge. Then the cartridge was sequentially washed with 1 mL water, 1 mL 5% MeOH. Elution was made into a labeled 12 \times 75 mm glass tube with 1 mL 100% MeOH. These eluates were further dried through a speed dryer at room temperature and reconstituted with 25% MeOH before analysis by HPLC with fluorescence detection.

Serum Sample Digestion Conditions

Diluted rat serum was used to determine the optimal conditions for enzymatic digestion and release of the AFB-Lys adduct such as incubation time, enzyme concentration used and temperature of incubation. A ratio of 1:4 (enzyme:total protein, w/w) was used for enzymatic digestion, as described previously²⁷. Pronase was prepared in PBS (10mg/mL, pH 7.4) and 250 μ L pronase was used to digest 150 μ L serum by incubation at 37°C water bath for 3 h, and the mixture was immediately put onto ice to stop the reaction until solid-phase extraction.

Serum Sample Preparation

Aliquots of 150 μ L human serum or diluted serum from AFB₁ treated rats were used for the method development. Solid-phase extraction with the Oasis Max™ cartridge was employed for the sample processing. The procedure for sample processing was same as that used in the Figure 3.1. The final eluates were dried under vacuum and reconstituted with 150 μ L of 25% MeOH, mixed thoroughly using multi-tube vortexer for 5 min, and centrifuged at 5000 rpm for 10 min at 4°C. Finally, 125 μ L of the supernatant was transferred into a HPLC vial for HPLC analysis.

HPLC Condition for AFB-Lys Analysis

The Agilent 1100 HPLC system with two mobile phases was used. Mobile phase A is 20 mM ammonium phosphate monobasic (pH 7.2) and B is 100% MeOH. A gradient was used for separation of AFB-Lys adduct and the run time was 25 min with a flow rate of 1.0 mL/min. Initial condition consisted of 70% A and 30% B and was

maintained for 5 min, then mobile phase A was linearly decreased to 35% in 15 min, retained for 3 min, then changed back to 70% in 2 min. The column was re-equilibrated for 5 min before the next run. A volume of 100 μL of the sample was injected into a reverse phase C18 column. The maximum excitation and emission wavelengths at 405 nm and 470 nm, respectively, were selected for the fluorescence detection.

Calibration Curve

Purified AFB-Lys was used to prepare standards which diluted in 25% MeOH for preparing calibration curves. Nine concentrations of AFB-Lys ranging from 10 to 5000 pg/mL were used to test the linearity on three separate days.

Limits of Detection

The known amount of AFB-Lys was added to human serum (20 pg/mg albumin) and then diluted with with normal human serum to determine the limit of detection (LOD). The limit of detection was defined as a S/N ratio of 2:1.

Accuracy and Precision

Accuracy was determined by diluting known amount of AFB-Lys with normal human serum to concentrations of 160 and 800 pg/mL. These diluted samples were analyzed with five replicates on three separate days. Intra-day precision was determined by the analysis of five replicates of low quality control (LQC, 160 pg/mL) and high quality control (HQC, 800 pg/mL) during a single run while inter-day precision was determined by analyzing five replicates of LQC and HQC for 3 separate analytical runs. The inaccuracy and imprecision were required to be within 15%.

Recovery Rate

Recovery rate was determined by adding known amounts of AFB-Lys adduct to normal human serum to produce triplicates at 50, 200 and 1000 pg/mL. These samples were incubated with 250 μ L pronase (as described above) at 37°C for 3 h. Sample preparation was done using the MAX™ cartridge as described above. After evaporation and reconstitution, the samples were analyzed in 25% MeOH. The peak area of standard without processing is represented as 100 percent. The recovery rate was calculated as the ratio between the mean peak areas of three processed samples and that of unprocessed standards.

Stability

Stability was determined using dilutions of serum obtained from AFB₁-treated rat. Room temperature stability up to 24 h was assessed by leaving the samples at room temperature for 12 h and 24 h followed by digestion and analysis. The stability of the AFB-Lys under conditions of repeated freezing and thawing was also examined for three cycles.

Serum Total Protein and Albumin Measurement

Total protein was measured before digestion of serum using a procedure modified from previous described method³³. Content of AFB-Lys adduct in serum was adjusted by albumin content. The albumin content in each sample was determined by a brom-cresol purple dye binding method, which measures a stable blue-purple color complex between dye and albumin with maximum absorbance at 600 nm⁴⁰.

Statistics and Graphics

Concentrations of AFB-Lys adduct were expressed as mean \pm SD and the ANOVA test was used to make comparisons among groups with SAS software (SAS Institute, Cary, NC, USA), followed by Tukey's test. *P*-values less than 0.05 were considered statistically significant.

Results

Quantification and Qualitative assessment of the Synthetic Standard

The synthetic standard was purified through fractional elution from the HPLC system and followed by evaporation and reconstitution in MeOH at 11.13 $\mu\text{g/mL}$ as determined by maximal UV absorbance at 400 nm. This standard is further confirmed through LC/MS and the mass chromatography as shown in Figure 3.2. The standard synthesized was qualitatively assessed using LC/MS. The SRM scan showed a transition from *m/z* of 454.5 and 456.5 (AFB-Lys adduct mol. wt 455.43²²), to a *m/z* of 394.2^{22,24}, which agreed with previously described data, confirming that the only detectable compound was authentic AFB-Lys adduct.

Binding Efficiency of Different Solid-Phase Extraction Cartridges

Commercially available solid-phase extraction cartridges were compared for their binding efficiency of AFB-Lys after enzyme digestion, shown in Table 3.1. HLB™ cartridge showed significantly lower binding efficacy of AFB-Lys adducts. While both MAX™ and Sep-Pak™ demonstrated similar binding efficacy, MAX™ cartridge was

preferred because it was less time-consuming and required less solvent compared with Sep-Pak™ cartridge.

Enzymatic Digestion of Aflatoxin Albumin Adduct from Serum

Several incubation periods were chosen and tested for release of AFB-Lys adduct using same amount of pronase (250 μ L). We found that the release of AFB-Lys adduct reached a peak at 3 h incubation, further elongation of incubation time did not produce higher amount (Figure 3.3), which was consistent with previous observations³⁶. The common procedure of inactivation of serum by incubation at 56°C for 30 min in water bath did not affect the release of AFB-Lys adduct from serum (data not shown here).

Calibration Curve Using Pure Synthetic AFB-Lys

Figure 3.4 shows the calibration curve of purified synthetic AFB-Lys adduct in 25% MeOH analyzed by HPLC separation with fluorescence detection. A total of nine concentrations from 10 to 5000 pg/mL were analyzed on three separate days. The correlation coefficient of the curve is 0.999. The linear range of this method encompasses the expected variance based on known human exposure levels.

LOD and LOQ

Human samples with known concentrations of AFB-Lys adduct were diluted to determine the limit of detection and the chromatograms are shown in Figure 3.5. We determined the LOQ to be 0.4 pg/mg albumin. The absolute LOD was 0.01 ng/mL, or 1 pg on-column (equivalent to 0.2 pg/mg albumin).

Accuracy and Precision

The accuracy of the assay based on three days ranged from 93.32% to 96.25% for LQC samples and from 92.23% to 97.25% for HQC samples, as shown in Table 3.2. The within day imprecision was determined to be 5.93% (n = 15) for LQC (mean = 0.16 ng/mL; 3.20 pg/mg albumin) and 2.87% (n = 15) for HQC (mean = 0.80 ng/mL, 16.0 pg/mg albumin) after solid-phase extraction and reconstitution in 25% MeOH.

Recovery Rate

Three different concentrations of AFB-Lys adduct were carried through the solid-phase extraction, evaporation and reconstitution process and the recovery rate was determined (Table 3.3). The recovery rates ranges from 72% to 94.44%. The mean \pm SD was $79.77 \pm 12.71\%$ at these three concentrations.

Stability

The stability of AFB-Lys was tested for up to 24 h at room temperature and 3 freeze-thaw cycles. Table 3.4 shows that there was no significant degradation of AFB-Lys adduct in diluted rat serum at 0.16 and 0.8 ng/mL concentrations (LQC and HQC) up to 12 h, the only significant decrease was observed at 0.8 ng/mL at 24 h. Table 3.5 shows that the three freeze-thaw cycles did no significantly affect the concentration.

Typical Chromatograms of Standard and Extracted AFB-Lys Adduct

Representative chromatograms from the synthetic standard and extracts from human serum are shown in Figure 3.6.

Discussion

Aflatoxin exposure remains an important food safety issue in that the Food and Agriculture Organization (FAO) estimates that 25% of the world's food crops are affected by mycotoxins, particularly aflatoxins. The worldwide burden of human hepatocellular carcinoma caused by long-term dietary exposure to aflatoxin is a public health concern ⁴¹. However, less-costly and more accurate methods for measuring human AFB₁ exposure are not readily for use in undeveloped countries and antibody based methods often vary in their sensitivity, thus making the results unreliable. Therefore, the development of a rapid method that doesn't require expensive mass spectrometer or antibodies or radioactive materials will allow developing countries to assess AFB₁ exposure without the need for major assistance from more developed countries. The purpose of the present study was to develop and validate such a method.

It is apparent that MAXTM and Sep-PakTM cartridges are better than the HLBTM cartridge with regard to binding and recovery of AFB-Lys adduct in serum. However, we found that the standard procedure for the Sep-PakTM cartridge took more time and had slightly higher variation compared with that of MAXTM, thus the MAXTM cartridge was chosen for solid-phase extraction in this assay.

Inactivation of possible pathogens is a common procedure before analysis of biological samples and subjecting serum samples to 56°C water bath for 30 min was tested and found to have no effect on the content of AFB-Lys in serum (data not shown). Enzyme content for digestion of treated serum was fixed at a ratio of 1:4 (w:w,

enzyme:total protein) based on total protein content present in serum²⁷. Incubation in the 37°C water bath resulted in a peak release of AFB-Lys after 3 h which was adopted in this assay while 4 h was used in the IDMS method. Less serum (150 µL) was needed for analysis compared to 250 µL serum in the IDMS assay³⁶. Thus, in terms of enzymatic digestion and volume of serum required, the method we have developed is similar to other validated methods.

The limit of detection in human serum was determined to be 0.4 pg/mg albumin, comparable to the reported 0.5 pg/mg albumin in the immune-affinity column based HPLC-fluorescence method³⁴ as well as in the isotope-dilution LC/MS/MS method³⁶. The ELISA performed directly on intact serum had a LOD of 100 pg/mg albumin and after digestion, 5 pg/mg albumin³⁴. The RIA method was reported to have a LOD of about 0.5 pmol (equivalent to 228 pg)³³. Thus, the HPLC fluorescence method developed in this study compare favorably with other validated methods.

A previous immune-affinity column based HPLC method reported a recovery of $83 \pm 6\%$ ³⁵. The uncorrected recovery for the IDMS extraction procedure was reported as $78 \pm 6.4\%$ ³⁶. Our results showed that the recovery of AFB-Lys adduct ranged from 72-94.4% with mean \pm SD of $79.77 \pm 12.71\%$. Inaccuracy and imprecision were within a range of 10% and thus fulfilled the requirements for an acceptable analytical method⁴², based on FDA guidelines.

Human sera from Texas subjects who participated in a collaborative project were analyzed using our method. The chromatograms displayed good separation of AFB-Lys

while commercial normal human serum showed less background interference at the same retention time. Taken together, this method for quantifying AFB-Lys in serum is fast and sensitive, which is amenable for application in large-scale population studies.

Acknowledgements

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Table 3.1. Binding efficacy of different cartridges (n=3)

Concentration (ng/mL)	MAX	HLB	Sep-Pak
0.16	0.12±0.01	0.05±0.01*	0.11±0.00
0.8	0.58±0.01	0.31±0.00*	0.57±0.05

*: $P < 0.05$, compared with other cartridges at same concentration.

Table 3.2. Accuracy and precision rates using diluted rat serum (n=5)

	AFB-Lys (ng/mL)	Observed value (ng/mL)	Inaccuracy rate (%)	Imprecision rate (%)
Day 1	0.16*	0.15±0.01	3.75	5.55
	0.80	0.86±0.02	7.77	2.76
Day 2	0.16	0.16±0.01	4.48	8.60
	0.80	0.82±0.02	2.80	2.39
Day 3	0.16	0.15±0.01	6.68	3.65
	0.80	0.79±0.03	2.75	3.47

* 0.16 ng/mL and 0.80 ng/mL were the LQC and HQC samples, respectively.

Table 3.3. Recovery rate of AFB-Lys adduct (n=3)

AFB-Lys (ng/mL)	Peak area (LU)		Recovery rate (%)
	Processed	Unprocessed	
0.05	4.53±0.38	4.80±0.20	94.44
0.2	14.77±0.55	20.27±0.12	72.87
1	73.2±1.35	101.67±0.51	72

Table 3.4. RT stability of AFB-Lys in serum (n=3)

Concentration (ng/mL)	Time (hour)		
	0	12	24
0.16	0.16±0.00	0.15±0.01	0.15±0.00
0.8	0.80±0.01	0.78±0.01	0.71±0.01*

*: $P < 0.05$, compared with 0 h at same concentration.

Table 3.5. Frozen-thawed stability of AFB-Lys in serum (n=3)

Concentration (ng/mL)	Frozen-thawed cycles			
	0	1	2	3
0.16	0.16±0.01	0.16±0.01	0.16±0.01	0.15±0.01
0.8	0.80±0.02	0.78±0.02	0.78±0.02	0.79±0.02

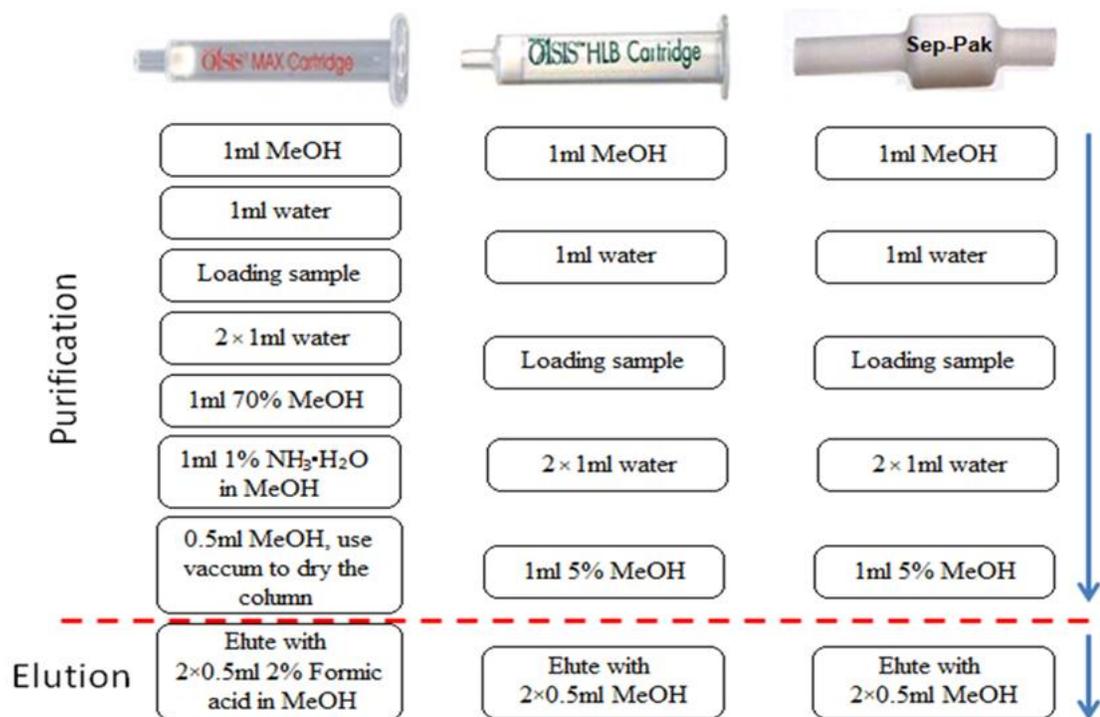


Figure 3.1. Flow chart of sample purification using different commercial cartridges.

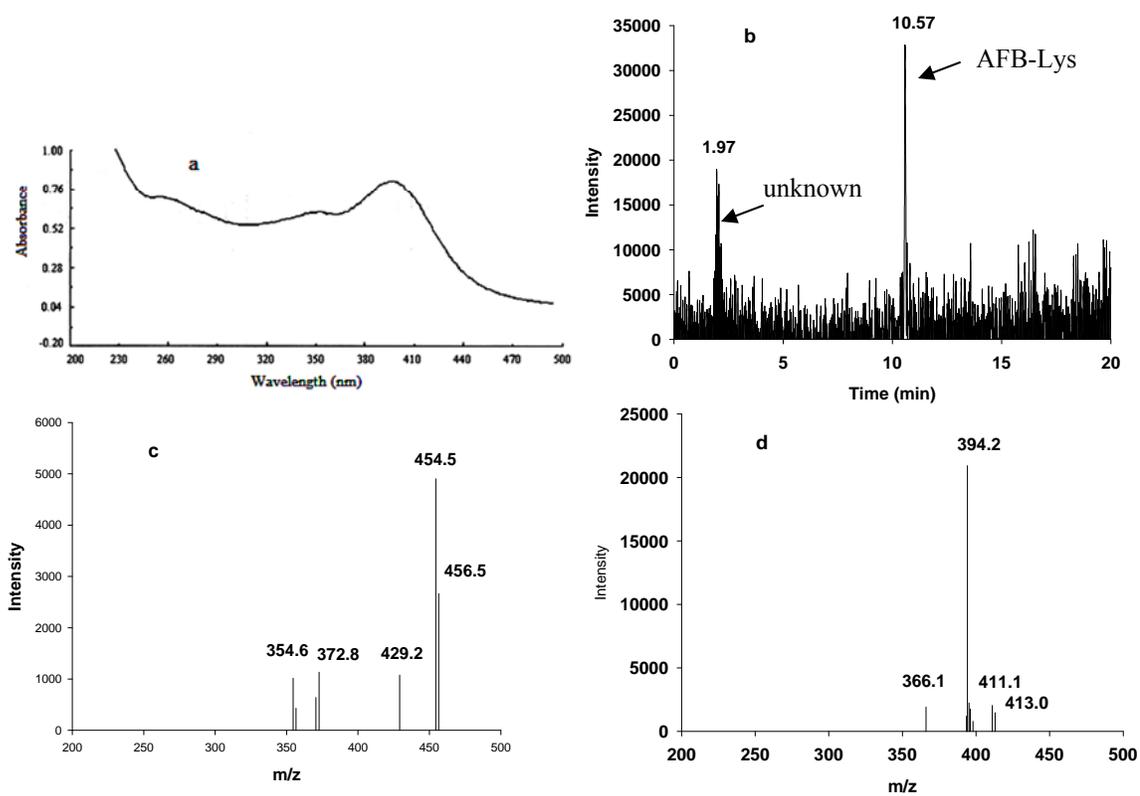


Figure 3.2. Chromatograms of synthetic AFB-Lys standard.

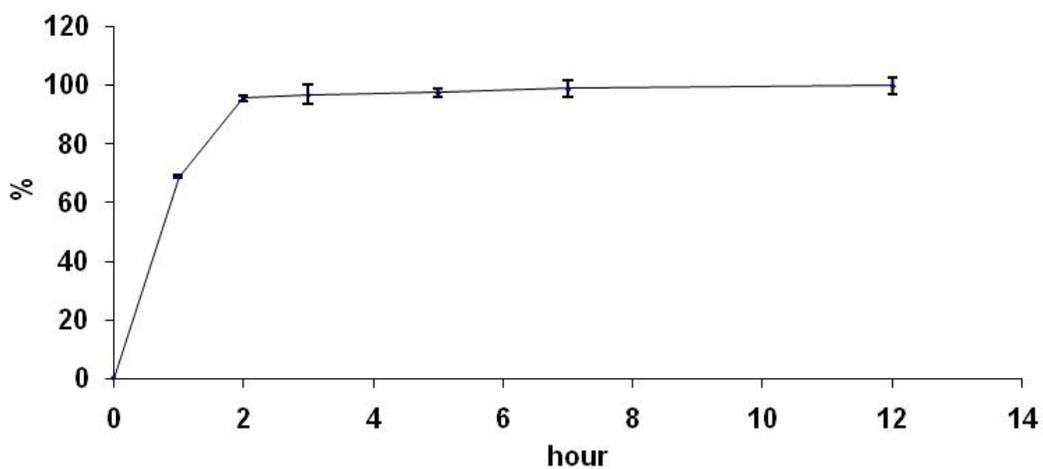


Figure 3.3. Effects of enzymatic incubation on the release of AFB-Lys from serum.

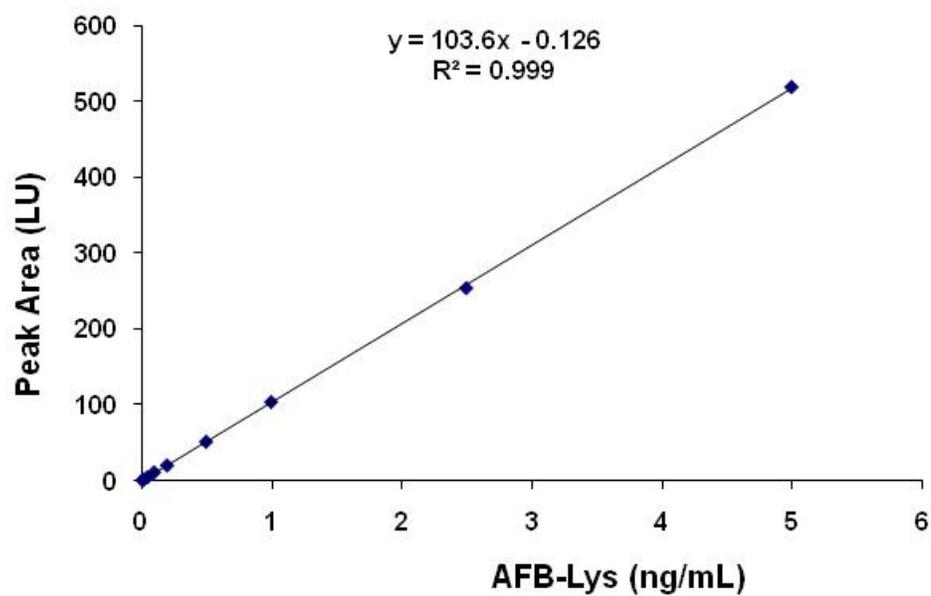


Figure 3.4. Calibration curve of AFB-Lys adduct.

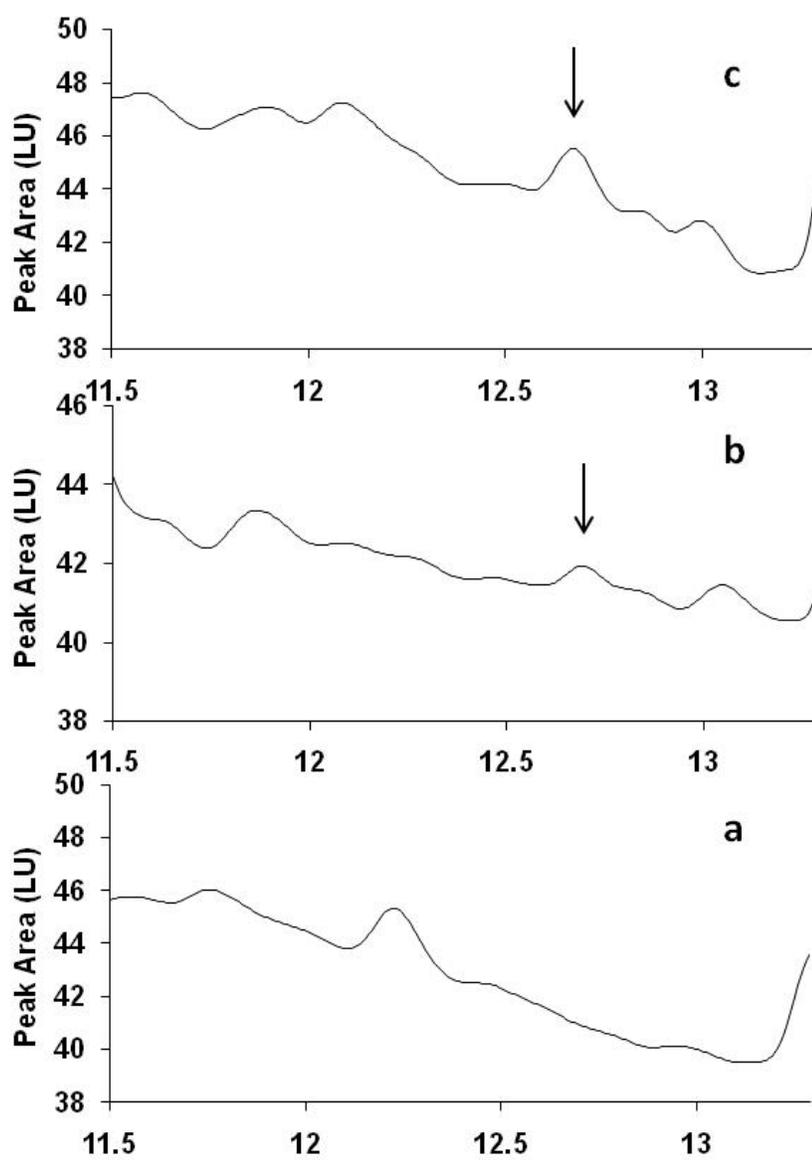


Figure 3.5. Detection limit of AFB-Lys adduct in human serum.

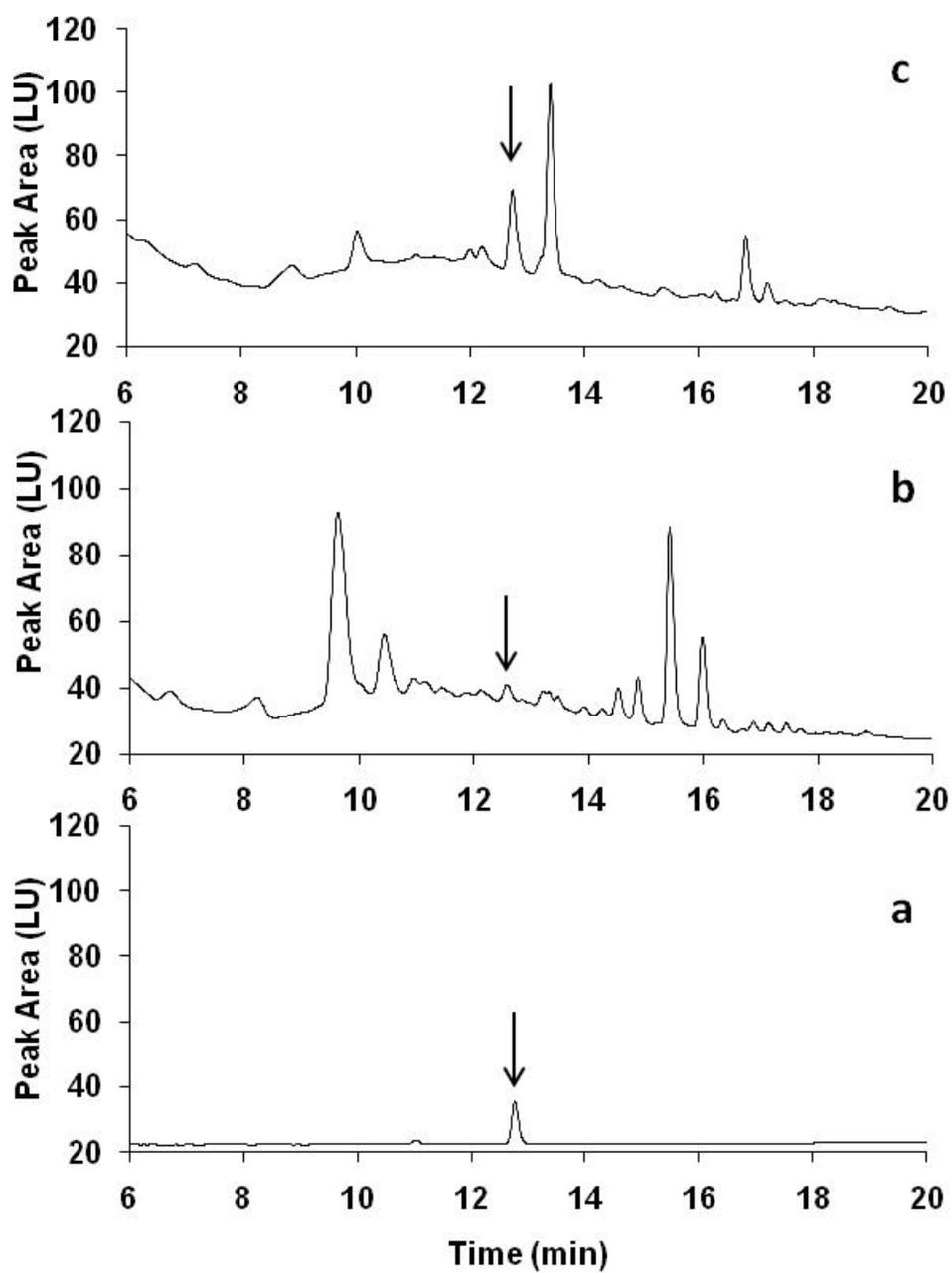


Figure 3.6. Typical chromatograms of synthetic and extracted AFB-Lys adduct

Figure Legends

Figure 3.1. Flow chart showing the steps used for sample purification using different commercial cartridges.

Figure 3.2. Purification of the AFB-Lys standard. a, UV/Vis spectrum of the AFB-Lys adduct showing maximal absorbance at 400 nm; b, chromatogram of the synthesized AFB-Lys adduct showing total ion current; c, mass spectra of the synthesized standard; d, collision-induced dissociation of the parent mass of 457 m/z scanning from m/z 200 to 500, the major fragment was 394.2 m/z, similar to that published by Scholl et al. (Reference #21).

Figure 3.3. Effects of length of enzymatic incubation on the release of AFB-Lys from serum.

Figure 3.4. Calibration curve of AFB-Lys adduct (10 to 5000 pg/mL) based on HPLC fluorescence detection as described in the methods.

Figure 3.5. Detection limit of AFB-Lys adduct in human serum. a), normal unexposed human serum; b), a processed sample containing 0.4 pg/mg albumin of AFB₁-Lys adduct; c) a processed sample containing 2.0 pg/mg albumin of AFB-Lys adduct. Retention time was about 12.7 min. Solid arrow indicated the peak for AFB-Lys adduct.

Figure 3.6. Typical chromatograms of synthetic and extracted AFB-Lys adduct a: 1 ng/mL calibration standard in 25% MeOH (equivalent to 20 pg/mg albumin); b: 1.5 pg/mg albumin AFB-Lys adduct in human serum; c: 30 pg/mg albumin in human serum. Retention time was about 12.7 min. Solid arrow indicated the peak for AFB-Lys adduct.

CHAPTER 4

PHYSIOLOGICALLY BASED TOXICOKINETICS OF SERUM AFLATOXIN

B₁-LYSINE ADDUCT IN F344 RATS ¹

¹ Qian, G. Tang, L. Wang, F. Guo, X. Massey, M.E. Williams, J.H. Phillips, T.D. and Wang, J.-S. 2012. *Toxicology*. In print. Reprinted here with permission of the publisher.

Abstract

Aflatoxin B₁-lysine adduct (AFB-Lys) is a reliable biomarker for aflatoxin exposure; however, a systematic toxicokinetic evaluation has not been reported. In this study, male F344 rats were orally exposed to single, or repeated, doses of AFB₁ and the toxicokinetics of serum AFB-Lys that followed treatments were investigated. A single-dose of AFB₁ increased serum AFB-Lys levels rapidly peaking at 4 h, followed by first-order elimination, through which the half-life was estimated to be 2.31 days. A physiologically based pharmacokinetic model showed that approximately 3.00-3.90% and 1.12-1.98% of the administered AFB₁ doses were converted to serum AFB-Lys adducts at 2 h and 24 h post treatment, respectively. Repeated AFB₁ exposure at 5-25 µg/kg body weight linearly increased serum AFB-Lys levels for 5 weeks in animals, resulting in a 1-1.5 times higher AFB-Lys level overall. This indicates the potential of this adduct as a reliable biomarker for repeated low dose exposure. Higher dose exposure at 75 µg/kg increased the level of AFB-Lys to a maximum at 2 weeks, followed by a gradual decrease to near plateau level up to 5 weeks. In conclusion, this study systematically evaluated the toxicokinetics of serum AFB-Lys adduct in F344 rats using a physiologically based pharmacokinetic model and robust statistical modeling analysis and provided a firm and clear understanding of the toxicokinetics of this biomarker.

Keywords: aflatoxin B₁; aflatoxin B₁-lysine adduct; biomarker; toxicokinetics.

Introduction

Aflatoxins (AF) are ubiquitous food contaminants produced mainly by *Aspergillus flavus* and *A. parasiticus*. Humans exposed to AF contaminated food are subject to a wide range of health effects, including acute poisoning (aflatoxicosis), hepatocellular carcinoma (HCC) and stunting in children¹⁻⁴. The uncovering of the etiological role of AF exposure is closely associated with the development and application of molecular markers relevant to AF exposure or risk^{5,6}.

The toxicology of aflatoxin B₁ (AFB₁) has been detailed elsewhere⁷. AFB₁ is metabolically activated by liver cytochrome P450 (CYP450) 1A2 and 3A4 to AFB₁-8,9-epoxide which can readily bind DNA to form AFB₁-N⁷-guanine adduct⁸ or react with albumin to produce AFB₁-albumin adduct if not conjugated with glutathione⁹. It has been shown that albumin is the only protein in serum that binds AFB₁ to a significant level¹⁰. Lysine is the primary amino acid group for the covalent binding of AFB₁ to albumin. Under enzyme digestion, AFB₁-lysine (AFB-Lys) is released from the adducted albumin⁹. This biomarker has been widely used to indicate human exposure status^{11, 12} and to assess human chemoprevention strategies^{13, 14}. Several previous studies have reported the partial toxicokinetics of the AFB-Lys adduct in animal models^{9, 15-17}. However, a complete study design relevant to human exposure (route, dose and duration) is lacking, and further studies are warranted to demonstrate the physiologically based toxicokinetics of the serum AFB-Lys biomarker.

Materials and Methods

Chemicals and Materials

Unlabelled aflatoxin B₁ (> 98% purity), dimethyl sulfoxide (DMSO), albumin determination reagent (bromocresol purple), and normal rat serum were purchased from Sigma Aldrich Chemical Co. (St. Louis, MO). Pronase (25 kU, Nuclease-free) was purchased from Calbiochem (La Jolla, CA). Protein assay dye reagent concentrate and protein standards were purchased from Bio-Rad Laboratories Inc. (Hercules, CA). Mixed mode solid phase extraction (SPE) cartridges were purchased from the Waters Corp. (Milford, MA). Authentic AFB-Lys was synthesized as previously described⁹. All other chemicals and solvents were of highest grade and purity available.

Animals

Male F344 rats (120-130 g) were purchased from Harlan Laboratories Inc. (Indianapolis, IN) and maintained under controlled conditions of temperature (22-25°C), humidity (50-70%), and light/dark cycles (12 h/12 h). Animals were fed with AIN-76A diet (Teklad, Madison, WI) and acclimated for one week before treatment. Feed and water were provided *ad libitum*. Animal use and care and the experimental protocol were approved by the Institutional Animal Care and Use Committee (IACUC) at the University of Georgia.

Experimental Design

For the single-dose study, rats (5 per group) were gavaged with 0, 50, 250, or 1,000 µg AFB₁/kg body weight (BW). AFB₁ was dissolved in DMSO with a gavage

volume of 50 $\mu\text{L}/100\text{g}$ BW. Animals were anesthetized and blood was collected by cardiac puncture at 2, 4, 8 h, and 1, 3, 5, 7, 14, and 21 days after treatment. For the repeated-dose study, rats (25 per group) were gavaged daily with 0, 5, 10, 25, or 75 $\mu\text{g}/\text{kg}$ BW for up to 5 weeks (5 days per week). Animals (5 per group) were anesthetized and blood samples were collected at weekly intervals and 24 h after the last dose. Serum samples were prepared and stored at -20°C until analysis.

Serum Sample Processing and Quantification of AFB-Lys Adducts

The method for serum processing and quantification of AFB-Lys adducts has been reported previously¹⁸. Albumin and total protein levels in serum were determined as previously described¹¹. Aliquots of each serum sample (150 μL) were digested by pronase (pronase:total protein, 1:4, w:w) at 37°C for 3 h to release AFB-Lys adducts. The digests were purified using Waters MAX SPE cartridges. After priming with methanol and equilibration with water, the loaded cartridge was sequentially washed by water, 70% methanol, and 1% ammonium hydroxide in methanol at a flow rate of 1mL/min. The eluate in 2% formic acid in methanol was vacuum-dried with a Labconco Centrivap concentrator (Kansas City, MO) and reconstituted for HPLC-fluorescence detection. The average recovery rate was 90%.

The Agilent 1200 HPLC-fluorescence system (Santa Clara, CA) was used for quantification of the AFB-Lys adducts. The mobile phases consisted of buffer A (20 mM $\text{NH}_4\text{H}_2\text{PO}_4$, pH 7.2) and buffer B (100% Methanol). The Zorbax Eclipse XDB-C18 reverse phase column (5 micron, 4.6×250 mm) was used and 100 μL was injected at a

flow rate of 1 mL/min. A gradient was generated to separate the AFB-Lys adduct within 25 min and the retention time of AFB-Lys was equal to approximately 12.3 min. AFB-Lys adduct was detected by fluorescence at the excitation and emission wavelengths of 405 nm and 470 nm, respectively. Quality assurance and quality control procedures were taken during analyses, which included simultaneous analysis of one authentic standard and a quality control sample daily. The limit of detection was 0.4 pg/mg albumin. The serum AFB-Lys level of each sample was adjusted by its albumin content accordingly.

Statistical Analysis

The curves of serum AFB-Lys formation and elimination after a single dose were fitted with zero-, first-, and second-order kinetic functions, which were used to determine the appropriate model for AFB-Lys kinetics. The lowest-order model with acceptable residuals ($r^2 > 0.90$) was considered to optimally describe the elimination curve. All modeling and residual determinations were performed with Sigma Plot 10.0 software (San Jose, CA) using the least sum of squares method. The half-life was computed from the fitting of the model selected.

Results

Single Dose Study

Following a single-dose of AFB₁, the levels of AFB-Lys in serum rapidly increased at 2 h and reached a peak at 4 h, followed by a non-linear decrease (Figure 4.1). Linear dose-response relationships were found between AFB₁ doses and serum AFB-Lys levels at both 2 and 24 h post treatment (Figure 4.2A and B, $r^2 = 0.98$ and 0.99 ,

respectively). In addition, the elimination of serum AFB-Lys followed first-order kinetics for both 50 ($r^2 = 0.9829$, $p < 0.001$) and 250 $\mu\text{g}/\text{kg}$ groups ($r^2 = 0.9745$, $p < 0.0001$) (Table 4.1). Non-linear decreases of serum AFB-Lys after 4 h were demonstrated based on model analysis for the 50 and 250 $\mu\text{g}/\text{kg}$ groups (Figure 4.3A and B). Significant linear decreases were also observed with the log (AFB-Lys) over time for the 50 and 250 $\mu\text{g}/\text{kg}$ groups. Half-lives of AFB-Lys were estimated to be 2.35 and 2.27 days from the curve fittings of the data from the 50 and 250 $\mu\text{g}/\text{kg}$ groups, respectively. The half-life was estimated based on the integrated slope (λ , as the regression coefficient of the fitted model) of the log (AFB-Lys) over time curves, i.e., $t_{1/2} = 0.693/\lambda$ (Figures 4.3C and D). The conversion ratios of AFB₁ to AFB-Lys following a single dose treatment were determined to be 3.00-3.90% at 2 h or 1.12-1.98% at 24 h post treatment (Tables 4.2 and 4.3), as estimated by a physiologically based pharmacokinetic model.

Repeated Dose Study

Repeated dosing with AFB₁ increased serum AFB-Lys for up to 5 weeks at all doses (Figure 4.4). A dose-dependent temporal increase in the AFB-Lys level was found in animals that received 5-25 $\mu\text{g}/\text{kg}$ AFB₁ over a 5-week study period. The level of AFB-Lys reached a maximum after 2 weeks, followed by a gradual decrease over time in the 75 $\mu\text{g}/\text{kg}$ group. To further examine the kinetics of the serum AFB-Lys after the repeated-dose treatment, we conducted different model analyses and found that the linear regression model best fit the data for the 5 (Figure 4.5A, $r^2 = 0.96$, $p < 0.01$), 10 (Figure 4.5B, $r^2 = 0.97$, $p < 0.01$) and 25 $\mu\text{g}/\text{kg}$ groups (Figure 4.5C, $r^2 = 0.98$, $p < 0.01$),

respectively. A Gaussian curve was the best fit for the 75 µg/kg group (Figure 4.5D, $r^2 = 0.86$, $p = 0.1634$). Linear dose-response relationships were also found between cumulative doses and serum AFB-Lys levels (Figure 4.6A, $r^2 = 0.99$, $p < 0.01$), as well as between the cumulative doses and the conversion ratio (AFB-Lys formation)/(AFB₁ dosed) after a 1-week treatment (Figure 4.6B, $r^2 = 0.99$, $p < 0.01$).

Discussion

In this study we evaluated the toxicokinetics of serum AFB-Lys adduct in F344 rats with special focus on doses relevant to acute (single) or chronic (repeated) human exposure. The peak concentration of serum AFB-Lys was found at 4 h in the current study. The mean AFB-Lys concentration of 2.56 ± 0.38 ng/mg albumin at 24 h after a single dose of 250 µg/kg was comparable to previously reported values in animals^{19, 20}. The highest dose of 1000 µg/kg was selected to simulate the extreme condition in which human acute aflatoxicosis occurred, as found in 2004-2005 outbreaks in Kenya, where a mean serum level of 3.2 ng/mg albumin for this biomarker has been reported in cases of aflatoxin poisoning before death¹².

Our physiological model based estimation of the conversion ratio of AFB₁ to AFB-Lys at 24 h after a single dose (1.12-1.98%) was similar to previously reported values of 1-3%^{9, 15}. The model generated half-life of serum AFB-Lys in this study (2.31 days, mean value of 2.35 and 2.27) was close to the previously reported values of 2.5 days⁹, 55 h¹⁵, and 2.6 days¹⁶ in rats. The elimination kinetics of serum AFB-Lys

followed a first-order kinetic decay model, as determined by comparisons of different model fittings.

It is worth noting that the dose range selected in the repeated experiment (5-75 $\mu\text{g}/\text{kg}$ BW) was relevant to the exposure levels in high-risk human populations, such as Kenya, Ghana, and Guangxi, China, where the mean AFB₁ levels were found to be 100-1000 $\mu\text{g}/\text{kg}$ in corn^{12, 21, 22}. We observed a linear increase of serum AFB-Lys for animals that received 5-25 μg AFB₁/kg daily, leading to a 1.0-1.5 times increase after 5 weeks compared to that after 1 week, which supports its potential use as a long-term biomarker. In addition, we observed that serum AFB-Lys reached a maximum after 2 weeks for animals that received a high daily dose of AFB₁ (75 $\mu\text{g}/\text{kg}$ BW). The pattern observed for the 75 $\mu\text{g}/\text{kg}$ group (Gaussian curve) may reflect a variation in the metabolic balance between AFB-epoxide formation and detoxification²³ or enzymatic induction of glutathione S transferase (GST)²⁴, and this coincided with the alternative signs of toxicity found at this dose.

It has to be pointed out that data for toxicokinetics presented in this study are different from previous studies in the following ways: (1) the current study employed a human relevant route of exposure (orally) rather than the i.p. injection adopted by Scholl et al¹⁶, which could better mimic the toxicokinetics of the biomarker *in vivo*; (2) we exposed animals with unlabelled AFB₁ rather than ³H-¹⁵ or ¹⁴C- labeled AFB₁⁹; (3) we used a physiologically based model to estimate the conversion of AFB₁ doses to serum AFB-Lys adducts rather than expression of the conversion ratio as AFB-Lys level (ng/mg

albumin)/AFB₁ dose ($\mu\text{g}/\text{kg}$ body weight). This conversion ratio as previously reported by Scholl et al. (2006) is not straightforward¹⁶ given the different units involved; (4) the current study used both single and repeated treatment procedures with relatively longer duration as compared to previous repeated treatment protocols of 14 days¹⁷ or 9 days¹⁶. One more highlight of this study is the use of robust statistical modeling analysis that greatly facilitates the understanding of the toxicokinetics of the AFB-Lys. This work also represents the first systematic evaluation of this biomarker using a complete design that incorporates relevant physiological basis into the toxicokinetic estimation. Data from this study provides a firm and clear understanding of the toxicokinetics of serum AFB-Lys adduct and will benefit future studies using this biomarker for linking to human health effects.

Conflict of Interest

The authors declare no conflict of interest.

Acknowledgements

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Table 4.1. Elimination kinetics of AFB-Lys adduct after a single dose.

Dose ($\mu\text{g}/\text{kg}$)	Zero order		First order		Second order	
	r^2	p -value	r^2	p -value	r^2	p -value
50	0.4913	0.0528	0.9829	<0.0001	0.7779	0.0038
250	0.4696	0.0607	0.9745	<0.0001	0.7534	0.0052

Zero order kinetics tested by linear regression of serum AFB-Lys overtime; first order kinetics tested by $\ln(\text{AFB-Lys})$ over time; second order kinetics tested by $1/\text{AFB-Lys}$ over time.

Table 4.2. Percentages of AFB₁ converted to serum AFB-Lys adduct 2 h after a single-dose treatment.

Dose ($\mu\text{g}/\text{kg}$)	Body weight (BW)	AFB ₁ administrated (μg)	AFB-Lys measured (nM/mL)	Estimated parameters			Conversion ratio (%)
				Whole blood volume (mL)	Whole serum volume (mL)	AFB ₁ dose in serum (nM)	
50	151.6	8.05	0.16	9.87	5.92	0.95	3.68
250	152.1	39.00	0.82	9.90	5.94	4.87	3.90
1000	150.8	148.00	2.42	9.82	5.89	14.25	3.00

Values are calculated from the average of 5 rats. Whole blood volume is calculated as $(\text{BW} \times 0.06 + 0.77)$, mL; whole serum volume is calculated as $(\text{whole blood volume}) \times 0.60$; AFB₁ dose in serum (nM) is calculated as $(\text{AFB-Lys measured, nM/mL}) \times (\text{whole serum volume, mL})$; conversion ratio is calculated as $(\text{AFB}_1 \text{ dose in serum, nM}) \times (\text{molecular weight of AFB}_1, 312\text{g/M}) / 1000 / (\text{AFB}_1 \text{ administrated, } \mu\text{g}) \times 100\%$. The calculation was based on assumptions that the binding of AFB₁ with proteins other than albumin is negligible in the blood stream and that serum AFB-Lys adducts are evenly distributed in the blood stream. Estimation of blood and serum volume was based on previous studies (Lee, H.B., Blaufox, M.D., 1985. *J. Nucl. Med.* 26, 72-76 and Everett, N.B., Simmons, B., Lasher, E.P., 1956. *Circ. Res.* 4, 419-424).

Table 4.3. Percentages of AFB₁ converted to serum AFB-Lys adduct 24 h after a single-dose treatment.

Dose ($\mu\text{g}/\text{kg}$)	Body weight (BW)	AFB ₁ administrated (μg)	AFB-Lys measured (nM/mL)	Estimated parameters			Conversion ratio (%)
				Whole blood volume (mL)	Whole serum volume (mL)	AFB ₁ dose in serum (nM)	
50	161.39	8.05	0.05	10.45	6.27	0.31	1.20
250	156.36	39.00	0.23	10.15	6.09	1.40	1.12
1000	148.38	148.00	1.62	9.67	5.80	9.40	1.98

Values are calculated from the average of 5 rats. Whole blood volume is calculated as $(\text{BW} \times 0.06 + 0.77)$, mL; whole serum volume is calculated as $(\text{whole blood volume}) \times 0.60$; AFB₁ dose in serum (nM) is calculated as $(\text{AFB-Lys measured, nM/mL}) \times (\text{whole serum volume, mL})$; conversion ratio is calculated as $(\text{AFB}_1 \text{ dose in serum, nM}) \times (\text{molecular weight of AFB}_1, 312\text{g/M}) / 1000 / (\text{AFB}_1 \text{ administrated, } \mu\text{g}) \times 100\%$. The calculation was based on assumptions that the binding of AFB₁ with proteins other than albumin is negligible in the blood stream and that serum AFB-Lys adducts are evenly distributed in the blood stream. Estimation of blood and serum volume was based on previous studies (*Lee, H.B., Blaufox, M.D., 1985. J. Nucl. Med. 26, 72-76* and *Everett, N.B., Simmons, B., Lasher, E.P., 1956. Circ. Res. 4, 419-42.*).

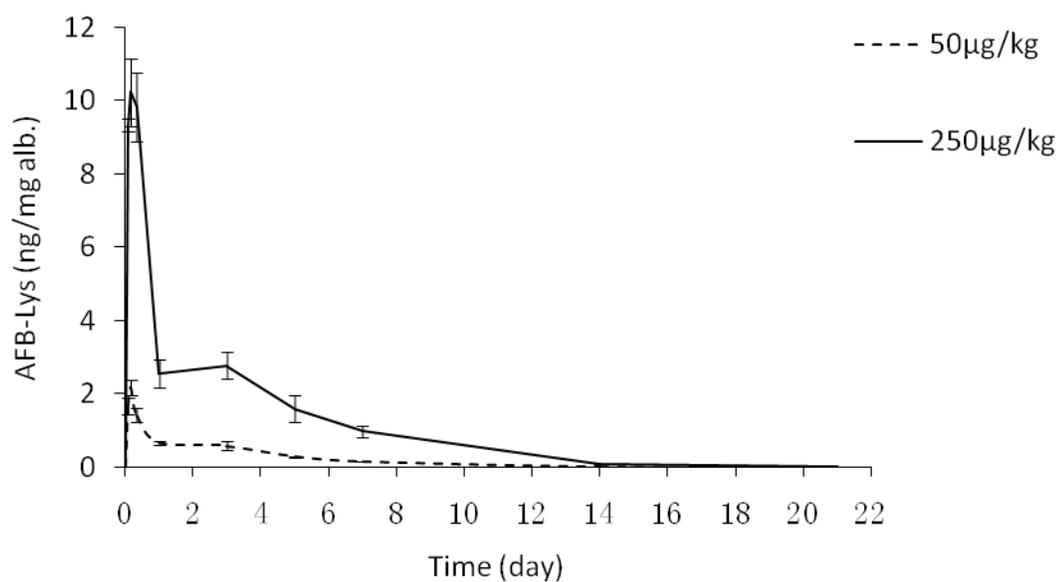


Figure 4.1. Serum concentrations of AFB-Lys adduct after a single-dose of AFB₁ exposure.

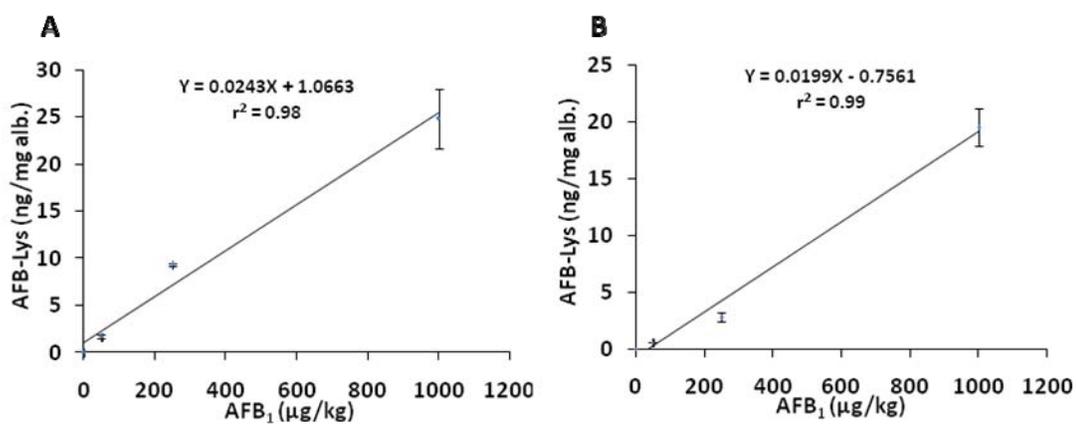


Figure 4.2. Linear relationships between AFB-Lys concentrations and AFB₁ doses after a single-dose treatment.

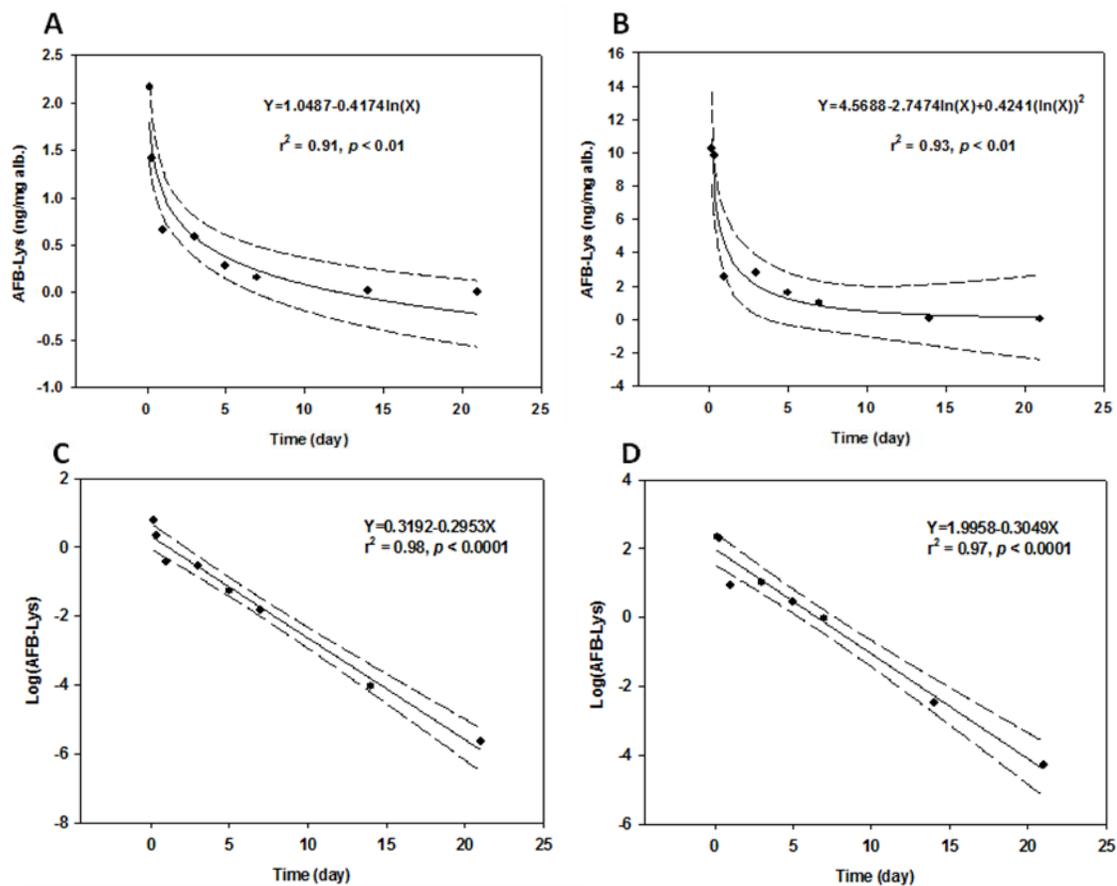


Figure 4.3. Elimination kinetics of serum AFB-Lys adduct after a single-dose of AFB₁ exposure.

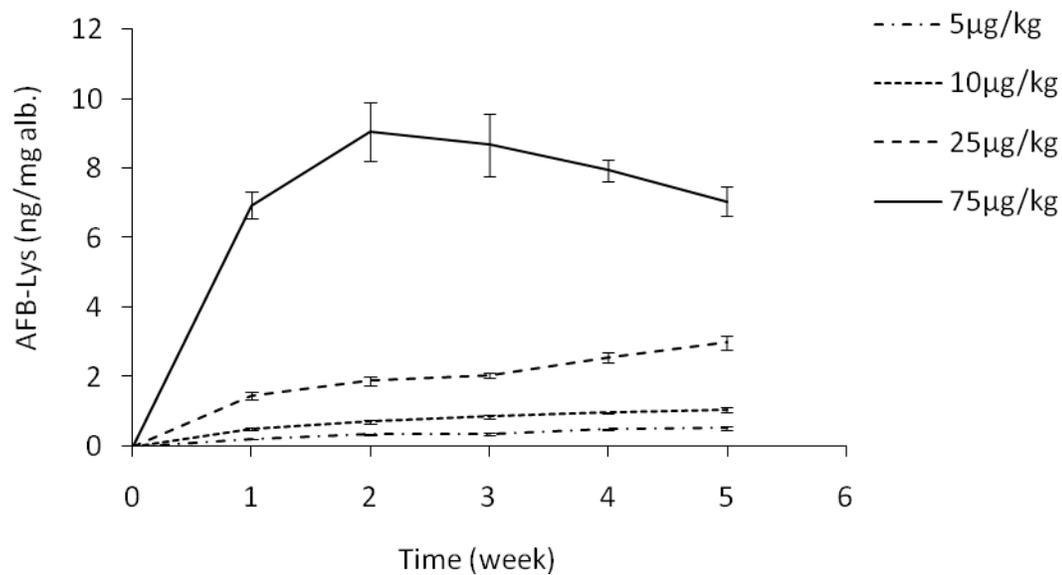


Figure 4.4. Serum concentrations of AFB-Lys adduct after repeated AFB₁ exposure.

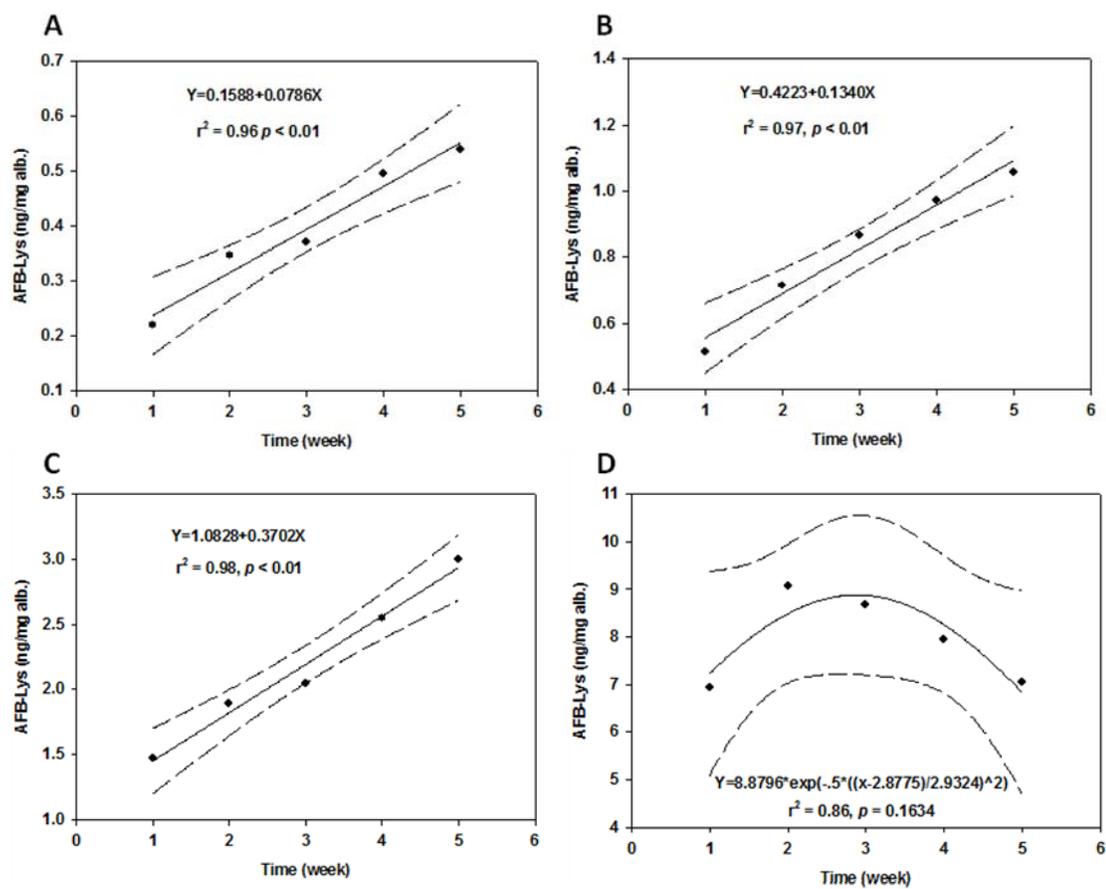


Figure 4.5. Regression analysis of serum concentrations of AFB-Lys adduct over repeated exposure.

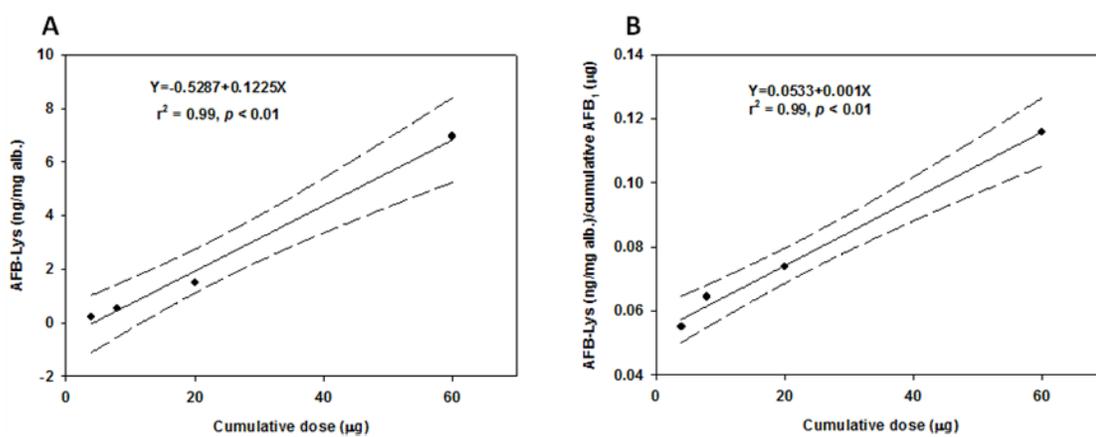


Figure 4.6. Dose-response and yield of AFB-Lys over cumulative AFB₁ doses.

Figure Legends

Figure 4.1. Serum concentrations of AFB-Lys adduct after a single-dose of AFB₁ exposure. The inset is an enlarged view of 0 to 24 h. Values shown are mean \pm SD (n=5).

Figure 4.2. Linear relationships between AFB-Lys concentrations and AFB₁ doses after a single-dose treatment. (A) 2 h ($r^2 = 0.98$) and (B) 24 h ($r^2 = 0.99$).

Figure 4.3. Elimination kinetics of serum AFB-Lys adduct after a single-dose of AFB₁ exposure. Nonlinear curve fitting of AFB-Lys over time was tested for the 50 (A) and 250 $\mu\text{g}/\text{kg}$ (B) groups. Linear curve fitting of the log (AFB-Lys) over time was tested for 50 (C) and 250 $\mu\text{g}/\text{kg}$ (D) groups. The slopes generated from Figs. C and D were used for calculating the half-life for serum AFB-Lys. Dashed lines indicate 95% CI.

Figure 4.4. Serum concentrations of AFB-Lys adduct after repeated AFB₁ exposure. Values shown are mean \pm SD (n=5).

Figure 4.5. Regression analysis of serum concentrations of AFB-Lys adduct over repeated exposure. (A), 5 $\mu\text{g}/\text{kg}$ group; (B), 10 $\mu\text{g}/\text{kg}$ group; (C), 25 $\mu\text{g}/\text{kg}$ group and (D), 75 $\mu\text{g}/\text{kg}$ group. Dashed lines indicate 95% CI.

Figure 4.6. Dose-response and yield of AFB-Lys over cumulative AFB₁ doses. Linear dose-response relationship between serum AFB-Lys concentrations and cumulative AFB₁ doses was fitted for 1 week after repeated-dose of AFB₁ ($r^2 = 0.99$, $p < 0.01$, A). Cumulative dose was calculated as dose ($\mu\text{g}/\text{kg}/\text{day}$) \times day \times body weight. Linear increase in yield of AFB-Lys over doses was also found ($r^2 = 0.99$, $p < 0.01$, B). Dashed lines indicate 95% CI.

CHAPTER 5
INTEGRATIVE TOXICOPATHOLOGICAL EVALUATION OF AFLATOXIN B₁
EXPOSURE IN F344 RATS ¹

¹ Qian, G. Wang, F. Tang, L. Massey, M.E. Mitchell, N.J. Su, J. Williams, J.H. Phillips, T.D. and Wang, J.-S. To be submitted to *Toxicologic Pathology*.

Abstract

In this study male F344 rats were orally exposed to a single-dose of aflatoxin B₁ (AFB₁) at 0, 50, 250 or 1000 µg/kg body weight (BW) or repeated-dose of 0, 5, 10, 25 or 75 µg/kg BW for up to 5 weeks. Biochemical and histological changes were assessed together with formation of AFB₁-lysine adduct (AFB-Lys) and liver foci positive for placental form glutathione S transferase (GST-P⁺). In single-dose protocol, serum AST, ALT and ALP were dose-related elevated with maximal changes (> 100-fold) at 3-day after treatment. Animals, received 250 µg/kg AFB₁, showed concurrent bile duct proliferation, necrosis and GST-P⁺ hepatocytes at 3-day, followed by liver GST-P⁺ foci appearance at 1-week. In repeated-dose protocol, bile duct proliferation and liver GST-P⁺ foci co-occurred after 3-week exposure to 75 µg/kg AFB₁, followed by proliferation foci formation after 4-week and dramatic ALT, AST and CK elevations after 5-week. Liver GST-P⁺ foci were temporally and dose-related induced. Serum AFB-Lys increased temporally at low doses (5-25 µg/kg), and reached maximum after 2-week exposure at 75 µg/kg. This integrative study demonstrate that liver GST-P⁺ cells and foci are sensitive biomarkers for AFB₁ toxic effect and correlated with bile duct proliferation and biochemical alterations in F344 rats.

Keywords: Aflatoxin B₁; aflatoxin B₁-lysine; placental form glutathione S transferase; histology; biochemistry.

Introduction

Aflatoxins (AF), especially aflatoxin B₁ (AFB₁), are common food contaminants in grains and groundnuts which represent a major group of harmful mycotoxins. The contamination of food by AF remains a significant public health problem worldwide, especially in the developing world where AF exposure has been reported to be associated with approximately 40% of all disease burden¹. The acute poisoning of AF (aflatoxicosis) in Kenya (2004 and 2005) had caused 125 deaths and more cases which resulted from the ingestion of heavily contaminated maize and maize products^{2, 3}. The adverse effects of AFB₁ on growth and development of children have become an increasing health concern^{4, 5}.

The histopathological effects and biochemical changes induced by AFB₁ exposure have been extensively studied in rat⁶⁻⁸ and chick^{9, 10}. However, only a few studies^{11, 12} reported the sequential biochemical and histological changes, in which rats were given either one single large dose of 3 mg/kg body weight (BW) or 7 mg/kg BW. In particular, no studies have been done to explore the correlations between biochemical and histopathological alterations with specific molecular targets via single or repeated treatment protocol using human relevant doses. Such data could be useful for delineation of functional and microstructural changes in liver and for evaluation of various intervention strategies to diminish the acute and chronic health effects due to AFB₁ exposure.

AFB₁ is metabolically activated by cytochrome P450 enzymes in liver to its reactive metabolite, AFB₁-8,9-epoxide, which binds to macromolecules resulting in DNA

adduct formation and carcinogenesis ¹³. AFB₁-8,9-epoxide can further be converted to AFB₁-8,9-diol that specifically binds to lysine in albumin and forms AFB₁-lysine adducts (AFB-Lys), which has been validated as a biomarker for indicating human exposure ¹⁴. This adduct and other AFB₁ biomarkers (e.g., AFB₁-N⁷-guanine adduct) have become critical tools for evaluation of chemopreventive agents in animals and humans ¹⁵.

Glutathione S transferase (GST) is a large family of metabolic enzymes comprised of several members, such as alpha, mu, pi, theta and others ¹⁶. GST in the alpha and pi classes is abundant in rat livers and inducible by drugs. Because of the putative role in detoxifying carcinogens these two isoenzymes have been the molecular target for chemoprevention ^{17, 18}. The pi form of GST was initially found in preneoplastic hepatocytes; in rat it was named GST-placental form (GST-P) as this form is the only form abundant in rat placenta (Ito et al., 1992). Rat liver GST-P positive (GST-P⁺) foci are a specific and reliable preneoplastic marker that have been widely used in animal carcinogenesis models ^{19, 20}. AFB₁ has been shown to induce GST-P⁺ hepatocytes or foci in rat liver both independently and synergistically with other toxins, such as fumonisin and microcystin LR ²¹⁻²³. However, the sequential development of GST-P⁺ foci caused by AFB₁ as well as the correlations with biochemical and histological changes in animal models remains largely unknown.

In this study we take an integrative approach to sequentially assess the toxic effects of AFB₁ in F344 rats following both single and repeated treatment protocols. The aims of this study are to validate the time course of development of liver GST-P⁺ foci, to evaluate the potential correlations of this effect biomarker with other histological and

biochemical changes caused by AFB₁ exposure, and to provide toxicopathological information for future intervention studies.

Materials and Methods

Chemicals and Reagents

AFB₁ (\geq 98% purity), normal rat serum, and dimethyl sulfoxide (DMSO) were purchased from Sigma-Aldrich Inc (St. Louis, MO). AFB₁ was prepared in DMSO to a stock solution and was diluted freshly to use. Rabbit anti-glutathione S transferase pi antibody was purchased from Abbiotec (San Diego, CA). Vectastain Elite ABC Kit and diaminobenzidine (DAB) substrate kit for peroxidase were purchased from Vector Laboratories (Burlingame, CA). All solvents were of the highest grade commercially available.

Animals

Male F344 rats (100-120 g) were purchased from Harlan Laboratory (Indianapolis, IN). Animals were individually housed under controlled light/dark cycle (12 h/12 h) with temperature of 22 \pm 2°C and relative humidity of 50-70%. Purified A1H 76A diet and tap water were provided *ad libitum*. Animals were maintained on AIN 76A feed for 1 week before treatment. Animal husbandry and care, AFB₁ treatment, and biological samples processing were reviewed and approved by the Institutional Animal Care and Use Committee (IACUC) at the University of Georgia.

Experimental Design

For single dose treatment, a total of 140 male F344 rats were randomly divided into 4 groups, 0, 50, 250 and/or 1000 μ g AFB₁/kg BW. Groups of 5 rats at each group were gavaged a single dose of AFB₁ and sacrificed at 2 h, 1, 3, 5, 7, 14, and 21 days after

treatment. For repeated-dose treatment, a total of 125 male F344 rats were randomly divided into 5 groups: 0, 5, 10, 25 and 75 $\mu\text{g AFB}_1/\text{kg BW}$. Animals were gavaged 5 days per week and 5 rats from each group were sacrificed at weekly intervals 24 h after the last dose. Gavage volume was set at 50 $\mu\text{L}/100 \text{ g BW}$. Blood and liver tissues were collected at sacrifice for examination of serum biochemistry, liver histology, and the GST-P⁺ foci formation. Animals were euthanized by cardiac puncture and exsanguination under isoflurane anesthesia.

Serum Biochemistry

Serum samples were collected by centrifuging the whole blood at 3000 rpm for 20 min at 4°C and stored at -20°C before analysis for serum biochemistry. All analysis was done with a Roche Hitachi 912 Chemistry Analyzer (Roche Diagnostics, Indianapolis, IN). The serum parameters measured were total serum protein (g/dL), albumin (g/dL), globulin (g/dL), albumin:globulin ratio (A/G ratio), serum calcium (mg/dL), phosphorus (mg/dL), glucose (mg/dl), BUN (mg/dL), creatinine (mg/dL), total bilirubin (mg/dl), alkaline phosphatase (ALP, U/L), creatine kinase (CK, U/L), alanine transaminase (ALT, U/L), aspartate transaminase (AST, U/L), g-glutamyl transpeptidase (GGT, U/L), amylase (U/L), cholesterol (mg/dl), serum sodium (mmol/L), potassium (mmol/L), Na/K ratio, chloride (mmol/L).

Serum AFB-Lys Adduct Analysis

The method for analysis of serum AFB-Lys has been described previously²⁴. Briefly, aliquots of each serum sample (150 μL) was digested with pronase (pronase:total protein, 1:4, w:w) at 37°C for 3 h to release AFB-Lys. The digests were purified by Waters MAX cartridges which were preprimed with methanol and equilibrated with

water. The loaded cartridge was sequentially washed by water, 70% methanol, and 1% ammonium hydroxide in methanol at a flow rate of 1mL/min. The purified AFB-Lys adducts were eluted with 2% formic acid in methanol, dried in a Labconco Centrivap concentrator (Kansas City, MO) and reconstituted for HPLC-fluorescence detection. The adduct concentrations were adjusted by serum albumin content. The average recovery rate was 90% and detection limit was 0.4 pg/mg albumin.

Liver Histology

Liver tissues were always taken from the right portion of the median lobe. Liver tissue blocks of approximately $6 \times 6 \times 1.5$ mm were routinely fixed in 10% neutral buffer formalin, dehydrated and embedded. Liver sections were cut at 5 μ m and stained with hematoxylin and eosin for light microscopic examination. Photographs were taken on an Olympus XC30 microscope with an Olympus UC30 digital camera (Olympus America Inc., Center Valley, PA) linked to a Celsense image analysis system (Celsense Inc., Pittsburgh, PA).

Liver GST-P Staining

Liver paraffin sections prepared as above were used for GST-P staining by the avidin-biotin-peroxidase complex (ABC) method²⁵. Briefly, liver sections were routinely hydrated and antigen was retrieved by heat. Sections were sequentially treated with primary rabbit anti-GST-P antibody (1:800), secondary goat anti-rabbit IgG (1:200) and ABC. The staining was done by adding diaminobenzidine (DAB). These sections were finally counterstained with hematoxylin for light microscopic examination. A GST-P⁺ focus was defined as a cluster of ≥ 5 hepatocytes positively stained with brown color.

Areas of each section were measured and the counting of GST-P⁺ cells or foci was based on section area.

Statistical Analysis

Data on body weights, serum biochemical parameters and AFB-Lys adducts were expressed as mean \pm standard deviation (SD). Statistical comparisons of these parameters were conducted using one-way analysis of variance (ANOVA) followed by Tukey's multiple comparison procedures. The raw data of body weight were used for statistical analysis while the biochemical parameters were analyzed following logarithmic transformation because of their right skewed distribution and non-equal variance. Correlation analysis was done through SigmaPlot 10.0 (San Jose, CA). A *p*-value of less than 0.05 was considered statistically significant.

Results

Body Weight Changes

Animals that received 1000 $\mu\text{g}/\text{kg}$ AFB₁ showed decreased body weight starting from 1 day after treatment ($p < 0.05$), as compared to the control (Figure 5.1A). For the 250 $\mu\text{g}/\text{kg}$ group, animals showed significantly decreased body weight gain at 3 and 5 days after treatment ($p < 0.05$). A dose-dependent decrease of body weight gain was found for animals that received repeated-dose treatment, however, significant decrease was found only in the 75 $\mu\text{g}/\text{kg}$ group starting from the second week as compared to control group ($p < 0.05$, Figure 5.1B). Animals in the 1000 $\mu\text{g}/\text{kg}$ group died within 7 days and the liver tissues were examined for histopathological changes and no animals died in any other group.

Serum Biochemistry after Single-Dose Treatment

Dose-dependent increases of serum AST and ALT activities were found at 1 and 3 days after single-dose treatment and these two enzymes were approximately 2- to 7-fold higher at 250 µg/kg group and 20- to 100-fold higher at 1000 µg/kg group, as compared to control group ($p < 0.05$, Table 5.1). AST and ALT were decreased after 1-week at 50 and 250 µg/kg groups ($p < 0.05$); ALT rather than AST returned to normal after 3-week. ALP activities were also elevated at 1000 µg/kg group at 1 and 3 days after treatment ($p < 0.05$, Table 5.1) while at lower doses ALP activities were found to be decreased starting at 3 days after treatment ($p < 0.05$). Serum CK was significantly decreased at 50 µg/kg group and elevated at 1000 µg/kg group at 1 day after treatment, as compared to control group ($p < 0.05$, Table 5.1). Starting from 3 days after treatment, CK was decreased in a dose-related manner and significantly lower than control group ($p < 0.05$). In addition, a dramatic elevation of total bilirubin was found at 1000 µg/kg group at 3 days after treatment with a mean \pm SD value of 15.49 ± 0.26 mg/dL, compared to all other groups including controls that have < 0.1 mg/dL total bilirubin (Data not shown in Table 5.1). Amylase activities were dose-dependently decreased at 1 day after treatment, followed by gradual recovery to normal at 1 week after treatment. Total protein and albumin were decreased at 1 and 3 days after treatment in a dose-dependent manner, which returned to normal after 1 week. Serum glucose levels were decreased at high doses (≥ 250 µg/kg) but elevated at low dose (50 µg/kg) at 1 and 3 days after treatment. Other serum biochemical parameters were less affected and data are not shown here.

Serum Biochemistry after Repeated-Dose Treatment

There were no obvious changes in profiles of serum biochemical parameters in animals after 1- or 3-week treatment with AFB₁ (Data not shown). However, repeated-dose treatment with 75 µg/kg AFB₁ for 5-week significantly increased serum AST and ALT activities by 1.02 and 0.19-fold, respectively, as compared to control group ($p < 0.05$, Table 5.2). Serum CK activities were decreased at 25 µg/kg and were dramatically elevated (approximately 3-fold) at 75 µg/kg group after 5-week treatment ($p < 0.05$, Table 5.2). ALP activities were significantly decreased at 25 µg/kg group ($p < 0.05$). Amylase activities and phosphorus levels were significantly elevated at 75 µg/kg group ($p < 0.05$). Glucose levels were significantly elevated at low doses (5 and 25 µg/kg) but decreased at high dose (75 µg/kg, $p < 0.05$). Bilirubin levels were less than 0.1 mg/dL in all groups after 5-week treatment (data not shown).

Serum AFB-Lys Levels

The levels of serum AFB-Lys after single-dose treatment are shown in Table 5.3. Dose-dependent changes of serum AFB-Lys were present at all time points. A quick rise at 2 h followed by decrease was found after single-dose treatment. Repeated-dose treatment for 5-week increased the adduct levels by approximately 1-fold at low doses (5-25 µg/kg), however, at high dose (75 µg/kg), the levels of the adduct were increased from start to a peak after 2-week treatment, followed by slow decrease over the 5-week experimental period (Table 5.4).

Liver Histological Changes after Single-Dose Treatment

The histological changes in rat liver induced by AFB₁ occurred as a function of both dose and duration. Dramatic bile duct proliferation occurred at 250 and 1000 µg/kg

groups at 3 days after treatment; massive periportal necrosis foci with inflammatory cells infiltration, excessive red blood cells appearing around hepatocytes, and destruction of liver lobes occurred only at 1000 $\mu\text{g}/\text{kg}$ group (Figure 5.2). Animals died in the 1000 $\mu\text{g}/\text{kg}$ group showed more severe liver damage but these changes were not reported here. Bile duct proliferation and periportal necrosis persisted at 250 $\mu\text{g}/\text{kg}$ group throughout the 3-week experimental period; however, no bile duct proliferation was found at 50 $\mu\text{g}/\text{kg}$ group throughout the experimental period (data not shown).

Liver Histological Changes after Repeated-Dose Treatment

The major histological changes in rat liver after repeated-dose treatment with AFB₁ included bile duct proliferation, periportal necrosis, and proliferation foci formation (Figures 5.3 and 5.4). Bile duct proliferation first appeared at 75 $\mu\text{g}/\text{kg}$ group after 3-week treatment, which progressed throughout the 5-week experimental period (Figure 5.3B). Bile duct proliferation was also found at 25 $\mu\text{g}/\text{kg}$ group (Figure 5.3A) but not at lower doses after 5-week treatment (data not shown). Necrosis was present in the periportal zone at doses higher than 10 $\mu\text{g}/\text{kg}$ group (Figure 5.3C and D). Proliferation foci were found only at 75 $\mu\text{g}/\text{kg}$ group after 4- and 5-week treatment (Figure 5.4A and B).

Liver GST-P⁺ Foci Formation after Single-Dose Treatment

Liver GST-P⁺ cells and foci occurred in a dose- and time-dependent manner (Table 5.5 and Figure 5.5). Liver GST-P⁺ foci were not found at 50 $\mu\text{g}/\text{kg}$ group throughout the experimental period; however, single GST-P⁺ cells were found starting from 1 week after a single-dose treatment. GST-P⁺ cells appeared first at 3 days for the 250 $\mu\text{g}/\text{kg}$ group, followed by foci formation at 1-week after treatment. The GST-P⁺ foci

persisted throughout the experimental period of 3-week. Interestingly, the highest dose of AFB₁ (1000 µg/kg) did not induce the formation of GST-P⁺ foci at 1-, 3- or 5-days after treatment. The liver GST-P⁺ foci observed in this study consisted of clear cells.

Liver GST-P⁺ Foci Formation after Repeated-Dose Treatment

The development of liver GST-P⁺ foci after repeated-dose treatment with AFB₁ exhibited dose- and time-dependency (Table 5.6 and Figure 5.6). No liver GST-P⁺ cells or foci were found in the control group at any time points examined; liver GST-P⁺ cells appeared at 5 µg/kg group after 3-week treatment, at 10 µg/kg after 2-week treatment and at 25 and 75 µg/kg groups after 1-week treatment, respectively. Liver GST-P⁺ foci appeared at 10 µg/kg group after 5-week treatment and at 25 and 75 µg/kg groups after 3-week treatment. There were no liver GST-P⁺ foci found at 5 µg/kg group throughout the experimental period. A good correlation between serum AFB-Lys adduct levels and the numbers of GST-P⁺ cells ($r = 0.99$, $p = 0.01$) or foci ($r = 0.97$, $p = 0.03$) after 5-week treatment was also found.

Discussion

In this integrative study we demonstrate the time course of development of biochemical, histological, and immunohistological changes and the connectivity between acute liver toxicity and carcinogenicity in both single-dose and repeated-dose treatment protocols (Figure 5.7). This is the first animal study to investigate the time course of development of liver GST-P⁺ hepatocytes and foci caused by AFB₁ exposure. Liver AST and ALT activities are early and sensitive indicators of acute hepatocyte injury or necrosis, in this case injury caused by AFB₁, which preceded the appearance of bile duct proliferation and periportal necrosis. The occurrence of GST-P⁺ hepatocytes was

temporally correlated with bile duct proliferation after single-dose treatment, while liver GST-P⁺ foci formation was correlated with bile duct proliferation in rat liver upon repeated-dose exposure. Serum AFB-Lys adduct was demonstrated as a reliable exposure biomarker especially through repeated low dose exposure. The doses of AFB₁ selected in this study were relevant to human exposure levels, which were found in cases of acute poisoning (such as in Kenya, up to 46.4 ppm AFB₁ in maize)³ and high-risk populations with chronic dietary exposure in China (estimated AFB₁ daily intake of 48.4 µg/male and 77.4 µg/female)²⁶. . An estimation of AFB₁ exposure, based on a 70 kg adult consuming 2 kg maize containing 1 ppm AFB₁, would be $2 \text{ kg} \times 1 \text{ ppm}/70 \text{ kg} = 26.7 \text{ µg/kg body weight}$. This is comparable to the selected doses in our study.

The transit rise of serum AFB-Lys at 2 h was in agreement with the quick absorption and metabolism of AFB₁ in rat liver²⁷. Compared to other parameters (amylase, proteins and glucose), dramatic increases of serum enzymes (ALT, AST and ALP) were found at 3 days after single treatment. These changes were in agreement with the massive necrosis in liver as observed at 3 days after treatment. It has been reported that a transit elevation of ALT and AST occurs at 48 h after three i.p. doses of AFB₁ (150 µg/kg) in Sprague-Dawley rats²⁸. The discrepancy is likely due to differences in animal species, route of exposure or examination period. Serum activities of AST and ALT returned to control group levels at 3-week for animals in the 50 but not 250 µg/kg group, indicating dose-related yet slow recovery process. The decreased CK after 1-day could be related to reduced muscle mass, evidenced by lowered body weight, and such finding has been reported in patients with alcoholic liver disease²⁹.

Repeated-dose treatment elevated serum AST and CK activities only after 5-week treatment with 75 $\mu\text{g}/\text{kg}$ AFB₁, and these changes occurred after the appearance of bile duct proliferation, indicating that these parameters are not sensitive indicators for repeated low dose AFB₁ exposure. The finding that CK was decreased at low dose (≤ 25 $\mu\text{g}/\text{kg}$) and elevated at high dose (75 $\mu\text{g}/\text{kg}$) may suggest different modes of action. In contrary, serum glucose levels were elevated at low doses and decreased at high dose, a finding similar to that in single-dose treatment. These outcomes could reflect a balance of different modes of actions, such as the decrease of glucose-6-phosphatase (G6P) activity⁸ and the newly expressed G6P in proliferating bile duct-like cells³⁰.

Bile duct proliferation and periportal necrosis are characteristic of AFB₁-induced liver injuries^{31, 32}, these histological changes slowly reached a maximum in rat liver at 3 days after a single dose of 250 or 1000 $\mu\text{g}/\text{kg}$ AFB₁ in this study. Animal received the highest dose (1000 $\mu\text{g}/\text{kg}$ AFB₁) died within 7 days of treatment. The possible reason may involve the massive hemorrhagic necrosis and liver lobular dissolution that became irreversible. Bile duct proliferation was constantly found in animals received 250 $\mu\text{g}/\text{kg}$ AFB₁ after 1 week but not at lower dose, indicating dose-related induction of this alteration. Among the three types of bile duct proliferation (typical, atypical and "oval cell"), the proliferation observed here is restricted to the periportal zone and usually has a well-defined lumen, which are characteristic of "typical" proliferation³³. The distorted liver microstructure might cause biliary duct hypertension that has been thought as the initiating factor in a bile duct ligated rat model.³⁴ Arguably, the newly produced bile duct could also be derived from a metaplasia of the periportal hepatocytes or from the

liver stem cells (or oval cells)³⁵. Although many studies have reported AFB₁ induced bile duct proliferation, the exact mechanisms remain to be uncovered.

Similar histological changes were found in animals received repeated-dose of AFB₁ treatment, such as bile duct proliferation and necrosis. In contrast to the "typical" bile duct proliferation, repeated low dose treatment caused more of an "oval cell" type of proliferation. The proliferation induces the formation of disorganized tubular structures with a poorly defined duct lumen³³. Since no specific staining has been done to distinguish these cells (bile duct epithelium, oval cell), it is likely that different types of proliferation may co-exist. The clear cell foci caused by repeated-dose treatment were early preneoplastic changes, which might return to normal or progress into hepatocellular carcinoma (HCC). For instance, HCC was reported in rats received 0.015 ppm AFB₁ for a period of 68-80 weeks with altered foci appearing within a few weeks' treatment³¹.

Liver GST-P is a specific biomarker that is markedly and localized increased in preneoplastic foci in rat¹⁶. This enzyme can be specifically induced by a variety of carcinogens, such as diethylnitrosamine (DEN) and 2-acetylaminofluorene (2-AAF)^{36,37}. Both single hepatocytes heavily positive for GST-P and liver GST-P⁺ foci have been used as markers of preneoplastic lesions in short-term carcinogenesis assays^{19,38}. In this study single GST-P⁺ hepatocytes were induced as early as 3 days after a single dose of 250 µg/kg AFB₁, demonstrating the strong potency of AFB₁ in initiating rat hepatocytes. A previous study demonstrated an even earlier onset of GST-P⁺ hepatocytes at 48 h after an i.p. injection of 2 mg/kg AFB₁ to young Fischer rats²³, and the differences were likely due to variance in animal age, dose and route of exposure. Liver GST-P⁺ foci were present at 250 µg/kg but not 50 µg/kg groups, which was suggestive of a non-observed

effect level for liver GST-P⁺ foci formation. Results from other studies on DEN and n-nitrosomorpholine^{39,40} were consistent with this finding, and the value of GST-P⁺ foci in dose-dependent carcinogenic studies has been reviewed⁴¹. In contrast, liver GST-P⁺ hepatocytes or foci were not induced in animals received 1000 µg/kg AFB₁. One possibility for the negative finding is the overwhelming of signaling pathways responsible for expressing GST-P, such as transforming growth factor β (TGF-β)⁴² and mitochondrial oxidative stress⁴³ mediated signaling pathways.

The appearance of liver GST-P⁺ hepatocytes was correlated with bile duct proliferation at 3 days after 250 µg/kg AFB₁ treatment, suggesting a synchronized onset. It was reported that 4-nitroquinoline 1-oxide induced placental GST expression was also correlated with cellular proliferation in a rat tongue carcinogenesis model⁴⁴. Therefore, it appears that a temporal overlapping exists between the acute toxicity (bile duct proliferation in particular) and the preneoplastic changes (liver GST-P⁺ cells and foci) at a non-lethal dose of AFB₁ exposure. While increased biliary pressure has been proposed as the initiating factor in bile duct proliferation³⁴ as mentioned above, the mechanisms for GST-P expression might also involve epigenetic mechanisms⁴⁵. However, the exact mechanisms for these effects require more efforts to uncover. In addition, serum AFB-Lys adduct levels increased over the 5-week experimental period at low doses (≤ 25 µg/kg), evident of its value for indicating chronic exposure. Large dose exposure (75 µg/kg) produced a plateau of this adduct after 2-week which was consistent with the occurrence of bile duct proliferation and GST-P⁺ foci at 3-week and may imply a variation of metabolic balance between its formation and detoxification.

Wild et al. have revealed the good correlation between serum AFB₁-albumin adducts and AFB₁-DNA adducts in multiple species including rats⁴⁶. Their data provide evidence for the application of AFB₁-albumin adducts from a mechanistic standpoint of view. Our study highlights the correlation between AFB₁-lysine adduct levels and formation of preneoplastic lesions: a good correlation between serum AFB-Lys adduct levels and numbers of liver GST-P⁺ cells or foci has been found after 5-week repeated exposure. It needs to mention that liver GST-P⁺ foci formation and serum enzyme changes (e.g., ALT and AST) are not specific, although these changes are sensitive to and can be correlated with AFB₁ exposure.

In conclusion, we demonstrate AFB₁-induced time course of development of liver GST-P⁺ foci and the correlations between formation of liver GST-P⁺ foci and bile duct proliferation in F344 rats. Given the early formation and the temporal correlation of Liver GST-P⁺ foci with bile duct proliferation after single or repeated AFB₁ treatment in F344 rats (as shown in this study), liver GST-P⁺ foci may be used as an endpoint to evaluate the efficacy of interventions in animal experiments, e.g., we have been conducting an animal experiment to assess the efficacy of NovaSil clay in preventing or reducing the toxic effects of co-exposure to AFB₁ and FB₁ through a short-term carcinogenesis rat model. The simultaneous examination of serum AFB-Lys adducts and liver GST-P⁺ positive foci might be useful for evaluation of chemopreventive agents in animal experiments.

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TABLE 5.1.—Selected alterations in serum biochemical parameters after single-dose treatment with AFB₁.

Time (day)	Dose (µg/kg)	AST (U/L)	ALT (U/L)	ALP (U/L)	CK (U/L)	Amylase (U/L)	Total Protein (g/dL)	Albumin (g/dL)	Glucose (mg/dL)
1	0	128.00±1.41 ^a	54.50±0.71 ^a	403.50±0.71 ^a	829.50±31.82 ^a	2261.00±18.38 ^a	6.30±0.00 ^a	4.35±0.07 ^a	219.00±2.83 ^a
	50	119.50±0.71 ^a	48.00±2.00 ^a	413.00±4.24 ^a	720.00±8.49 ^b	2233.50±2.12 ^a	6.20±0.00 ^b	4.30±0.00 ^a	242.00±2.83 ^b
	250	274.00±7.07 ^b	174.50±7.78 ^b	398.00±8.48 ^a	828.00±2.83 ^a	1915.00±21.21 ^b	5.95±0.07 ^c	4.25±0.07 ^a	169.00±2.83 ^c
	1000	2489.50±102.53 ^c	1854.50±106.77 ^c	485.50±3.53 ^b	1122.50±6.36 ^c	488.50±10.61 ^c	5.25±0.07 ^d	3.90±0.00 ^a	93.00±1.41 ^d
3	0	107.00±4.24 ^a	49.50±3.54 ^a	339.50±10.60 ^a	638.50±17.68 ^a	2415.00±110.31 ^a	6.35±0.21 ^a	4.45±0.07 ^a	212.50±7.78 ^a
	50	97.50±2.12 ^a	48.00±1.41 ^a	280.00±5.66 ^b	485.00±9.90 ^b	2483.00±21.21 ^a	6.05±0.21 ^{a,b}	4.25±0.07 ^a	234.00±5.66 ^a
	250	442.00±1.41 ^b	355.00±1.41 ^b	298.50±0.71 ^b	569.50±6.36 ^c	2523.00±29.70 ^a	5.60±0.00 ^b	3.80±0.00 ^b	188.50±0.71 ^b
	1000	10232.00±284.26 ^c	6150.50±58.69 ^c	639.50±6.36 ^c	414.50±4.95 ^d	1002.00±14.14 ^b	3.75±0.07 ^c	3.00±0.00 ^c	60.00±1.41 ^c
7	0	143.00±1.41 ^a	65.00±3.00 ^a	362.50±2.12 ^a	851.50±2.12 ^a	2510.50±23.33	6.10±0.00	4.20±0.00	200.50±0.71
	50	87.50±4.95 ^b	38.00±2.83 ^b	251.50±10.61 ^b	505.50±16.26 ^b	2161.00±83.44	5.85±0.21	4.10±0.14	204.50±6.36
	250	81.00±7.07 ^b	42.50±0.71 ^b	250.00±11.31 ^b	383.50±51.62 ^b	2384.00±124.45	6.10±0.14	4.20±0.00	213.00±15.56
21	0	98.00±2.83 ^a	43.50±2.12	245.50±2.12 ^a	618.50±6.36 ^a	2663.00±14.14	6.35±0.07	4.30±0.00	182.00±1.41 ^a
	50	87.50±6.36 ^{a,b}	36.50±2.12	209.00±2.83 ^b	443.50±57.28 ^{a,b}	2340.50±23.33	6.25±0.07	4.35±0.07	198.00±12.73 ^{a,b}
	250	74.00±4.24 ^b	43.00±4.24	224.50±0.71 ^c	327.00±32.53 ^b	2636.00±141.42	6.35±0.07	4.40±0.00	224.00±9.90 ^b

Note: values are means ± SD for 5 rats. Values that do not share a label (a letter or blank) are significantly different ($p < 0.05$).

TABLE 5.2.—Selected alterations in serum biochemical parameters after 5-week treatment with AFB₁.

Dose (µg/kg)	AST (U/L)	ALT (U/L)	ALP (U/L)	CK (U/L)	Amylase (U/L)	Glucose (mg/dL)	Phosphorus (mg/dL)
0	123.00±1.41 ^a	57.00±0.71 ^a	212.50±7.78 ^{a,b}	872.00±31.11 ^a	2644.00±73.54 ^a	164.00±2.83 ^a	10.27±0.36 ^a
5	111.50±7.78 ^a	47.50±0.71 ^b	200.00±8.49 ^{b,c}	819.00±35.36 ^a	2817.00±98.99 ^{a,b}	202.50±7.78 ^{b,c}	10.90±0.46 ^a
10	119.50±4.95 ^a	52.00±1.41 ^{a,b}	195.00±8.49 ^{b,c}	797.50±36.06 ^a	2735.00±96.17 ^{a,b}	180.50±6.36 ^{a,b}	9.86±0.47 ^a
25	106.50±4.95 ^a	50.50±0.71 ^b	178.00±5.66 ^c	429.00±14.14 ^b	2676.00±82.02 ^a	212.50±7.78 ^c	9.62±0.21 ^a
75	248.00±8.49 ^b	68.00±2.83 ^c	234.50±9.19 ^a	2556.50±101.12 ^c	3150.50±142.13 ^{b,c}	87.00±2.83 ^d	20.28±0.93 ^b

Note: values are means ± SD for 5 rats. Values that do not share a label (a letter or blank) are significantly different ($p < 0.05$).

TABLE 5.3.—Serum concentrations of AFB-Lys adduct after single-dose treatment with AFB₁.

Dose (µg/kg)	AFB-Lys (ng/mg albumin)						
	2 h	1-day	3-day	5-day	1-week	2-week	3-week
0	ND	ND	ND	ND	ND	ND	ND
50	1.68±0.22	0.66±0.05	0.58±0.12	0.28±0.02	0.16±0.01	0.02±0.00	ND
250	9.34±0.18	2.56±0.38	2.78±0.37	1.59±0.37	0.98±0.16	0.08±0.01	0.01±0.00
1000	24.89±3.17	19.97±1.67	NA	NA	NA	NA	NA

Note: ND, not detectable; NA, not available.

TABLE 5.4.—Serum concentrations of AFB-Lys adduct after repeated-dose treatment with AFB₁.

Dose (µg/kg)	AFB-Lys (ng/mg albumin)				
	1-week	2-week	3-week	4-week	5-week
0	ND	ND	ND	ND	ND
5	0.22±0.01	0.35±0.02	0.37±0.04	0.50±0.03	0.54±0.05
10	0.51±0.03	0.71±0.07	0.87±0.06	0.97±0.03	1.06±0.08
25	1.48±0.12	1.89±0.12	2.05±0.08	2.55±0.15	3.00±0.20
75	6.94±0.38	9.06±0.84	8.67±0.90	7.96±0.31	7.06±0.43

Note: ND, not detectable.

TABLE 5.5.—Liver GST-P⁺ cells and foci induced by single-dose treatment with AFB₁.

GST-P ⁺	Dose (µg/kg)	Time (days)					
		1	3	5	7	14	21
Cells	50	-	-	2.47±3.50 (3/5)	10.51±5.17 (5/5)	6.62±9.36 (4/5)	5.98±0.98 (5/5)
	250	-	22.72±17.25 (5/5)	90.37±23.17 (5/5)	189.80±102.78 (5/5)	142.23±53.24 (5/5)	92.62±32.29 (5/5)
Foci	50	-	-	-	-	-	-
	250	-	-	-	4.22±4.10 (3/5)	6.18±6.01 (5/5)	5.62±6.43 (5/5)

Note: data are mean values from 5 liver sections, each from one individual rat (No./cm²). -, not found. Data in brackets are (No. of animals affected/No. of animals examined).

TABLE 5.6.—Liver GST-P⁺ cells and foci induced by repeated-dose treatment with AFB₁.

GST-P ⁺	Dose (µg/kg)	Time (week)				
		1	2	3	4	5
Cells	5	-	-	14.99±12.55 (3/5)	18.83±10.28 (2/5)	16.74±13.20 (2/5)
	10	-	7.66±0.99 (5/5)	12.79±11.11 (4/5)	37.88±17.78 (5/5)	61.35±51.88 (5/5)
	25	20.97± 7.46 (5/5)	19.42±5.98 (5/5)	30.02±14.59 (5/5)	56.60±16.28 (5/5)	95.35±34.91 (5/5)
	75	21.47±11.62 (5/5)	63.36±11.57 (5/5)	245.42±92.78 (5/5)	203.65±82.77 (5/5)	259.80±41.19 (5/5)
Foci	5	-	-	-	-	-
	10	-	-	-	-	1.49±2.57 (2/5)
	25	-	-	1.36±1.22 (3/5)	1.12±1.94 (2/5)	9.24±12.78 (3/5)
	75	-	-	3.01±2.87 (3/5)	39.09±4.67 (5/5)	75.99±23.58 (5/5)

Note: data are mean values from 5 liver sections, each from an individual rat (No./cm²). -, not found. Data in brackets are (No. of animals affected/No. of animals examined).

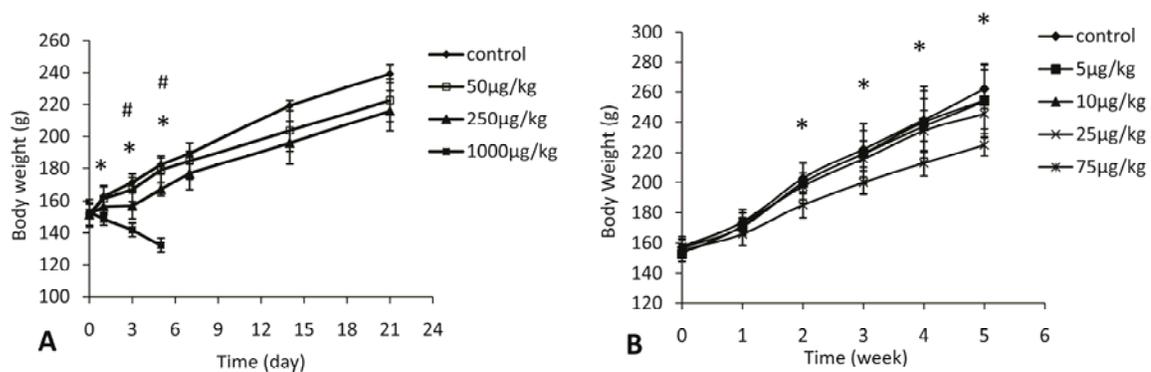


FIGURE 5.1.—Effects of AFB₁ on body weight gain after single-dose or repeated-dose treatment with AFB₁

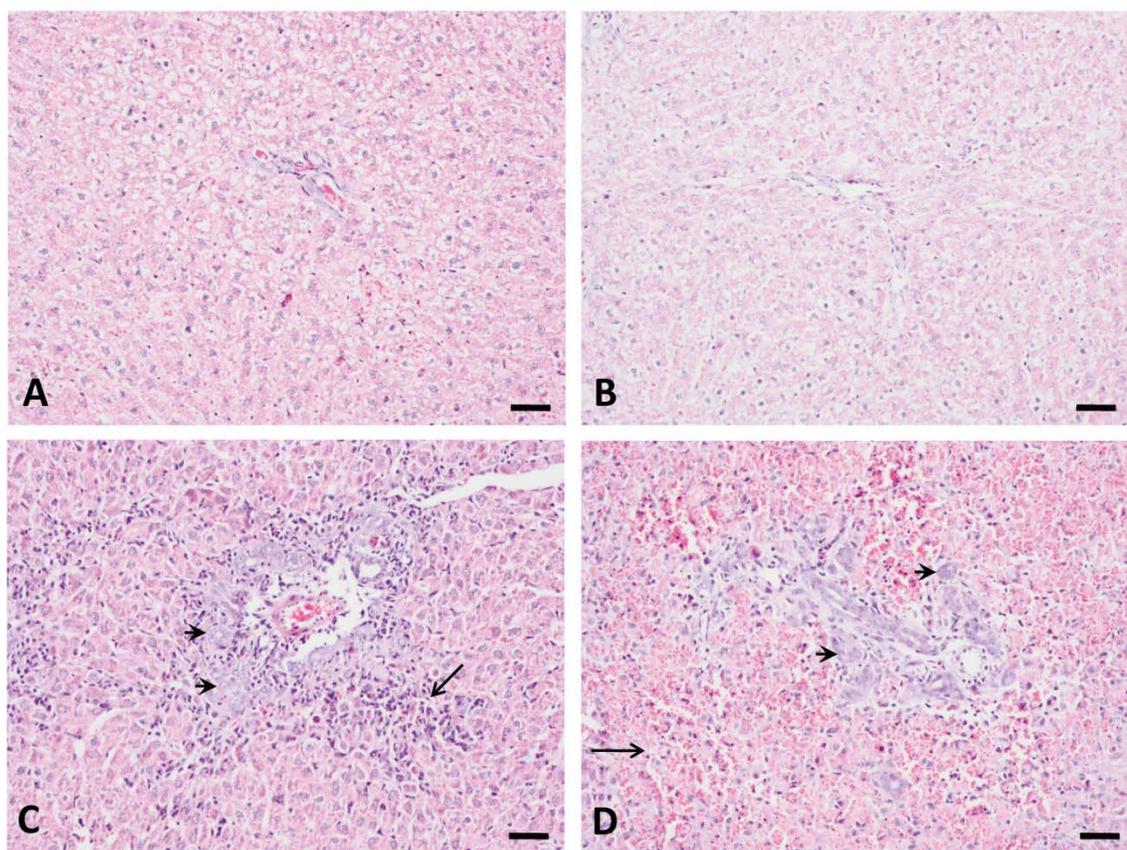


FIGURE 5.2.—Bile duct proliferation and periportal necrosis induced by AFB₁ at 3 days after single-dose treatment

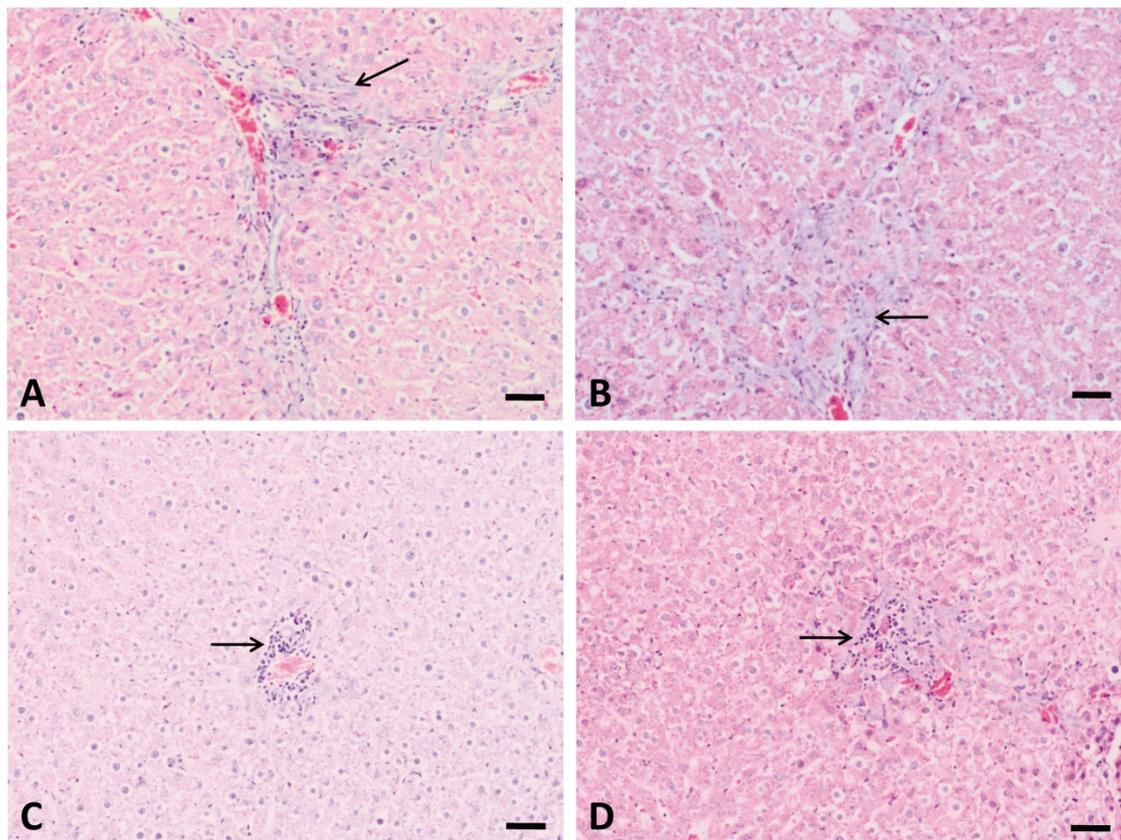


FIGURE 5.3.—Bile duct proliferation and periportal necrosis induced by 5-week treatment with AFB₁

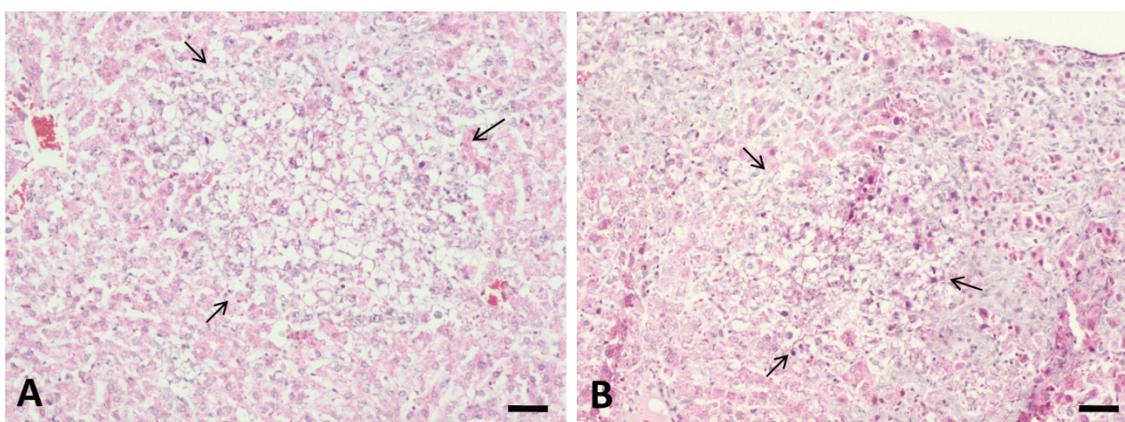


FIGURE 5.4.—AFB₁ induced proliferation foci in liver after repeated-dose treatment

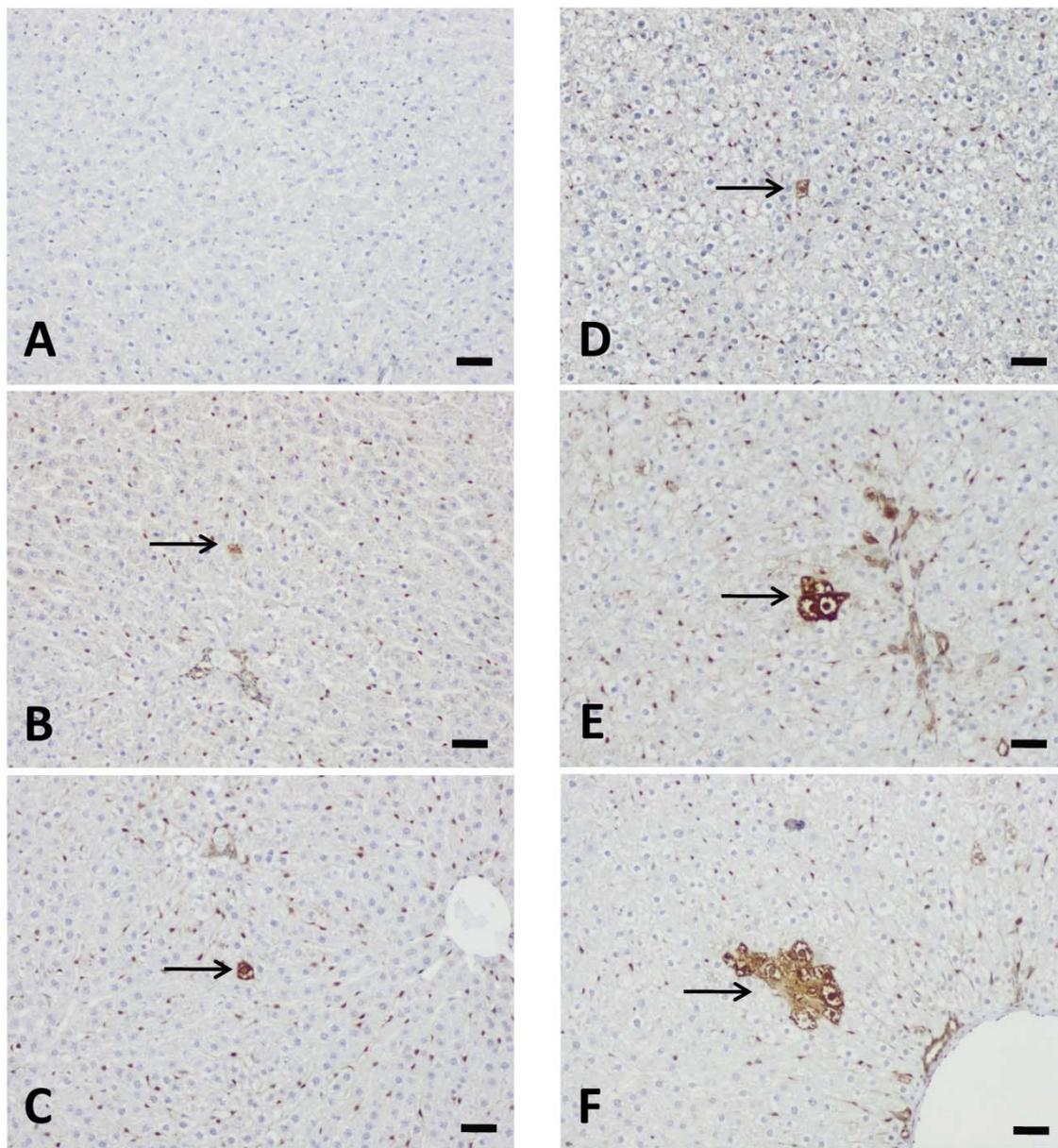


FIGURE 5.5.— AFB_1 induced liver GST-P⁺ cells or foci formation after single-dose treatment with AFB_1

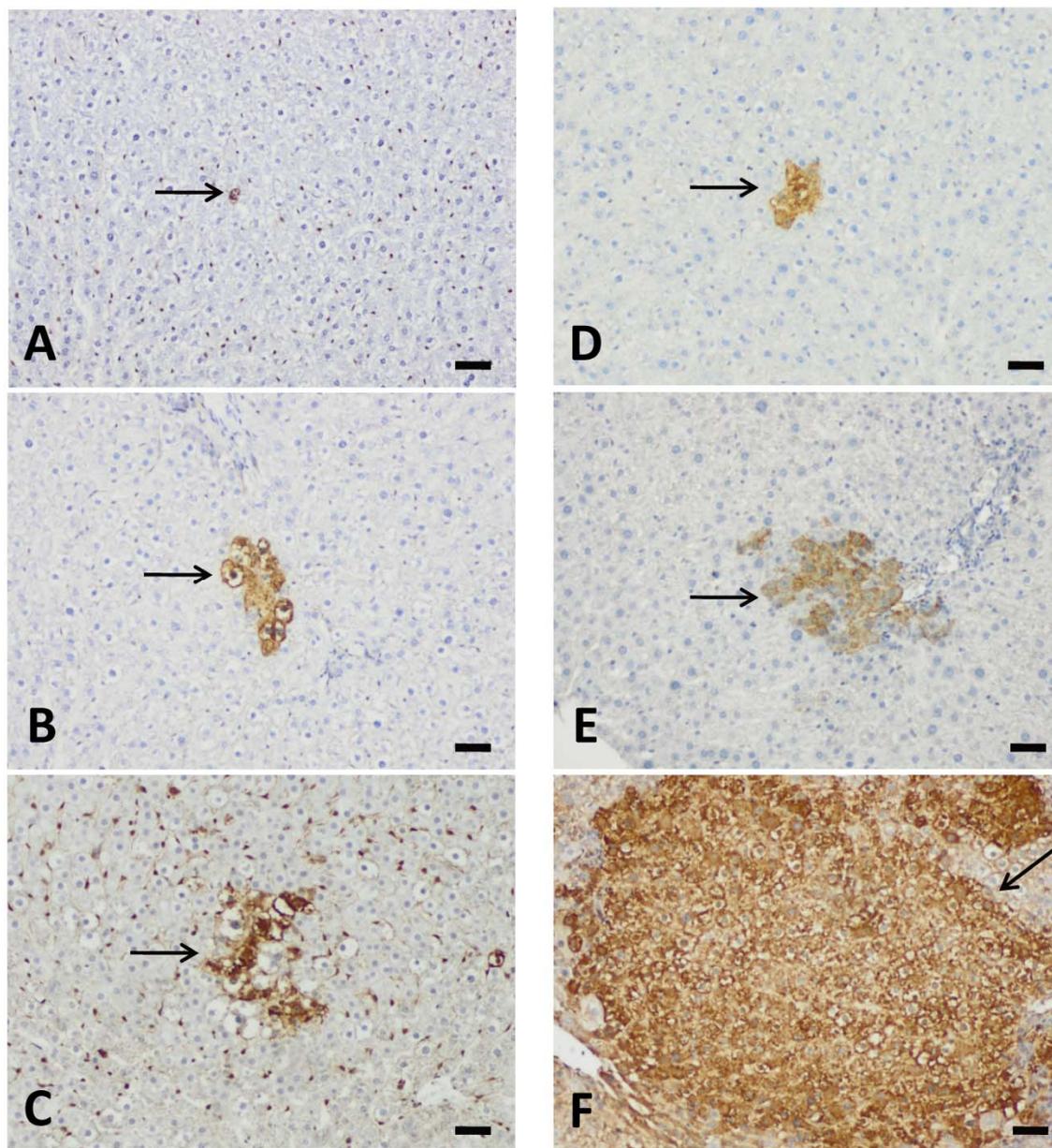


FIGURE 5.6.— AFB_1 induced liver GST-P⁺ cells or foci formation after repeated-dose treatment with AFB_1

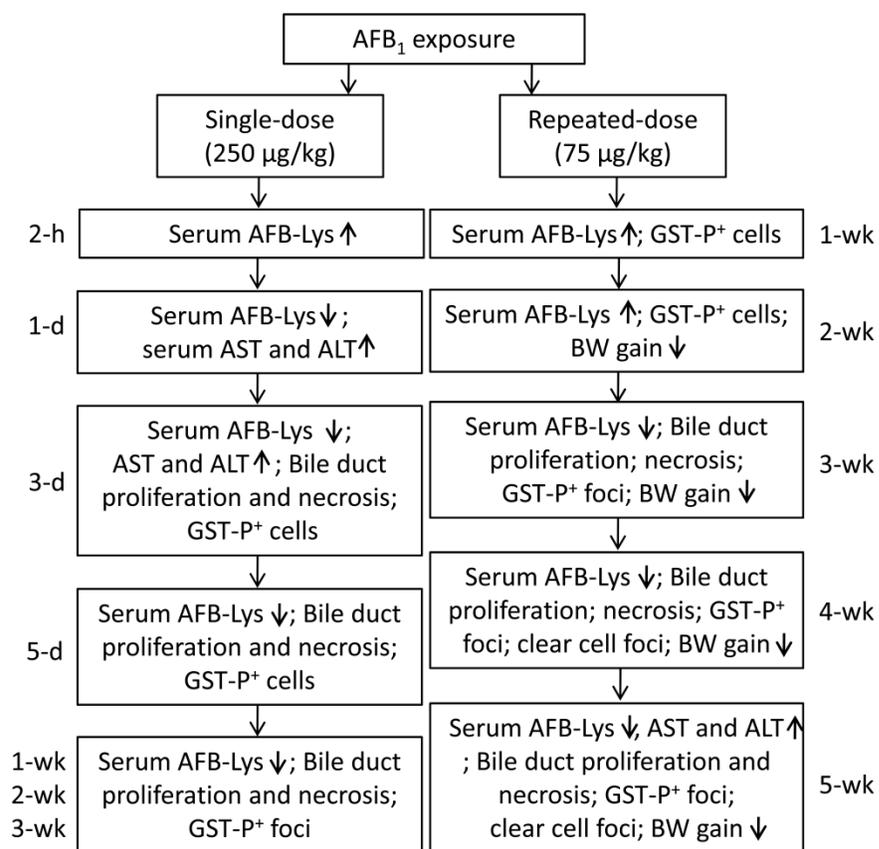


FIGURE 5.7.—Summary of the integrative toxicopathological findings in this study

Figure Legends

FIGURE 5.1.—Effects of AFB₁ on body weight gain after single-dose or repeated-dose treatment with AFB₁. *, $p < 0.05$, the highest dose group vs control group; #, $p < 0.05$, the highest dose group vs 50 $\mu\text{g}/\text{kg}$ group.

FIGURE 5.2.—Bile duct proliferation and periportal necrosis induced by AFB₁ at 3 days after single-dose treatment. No obvious histological changes occurred for control (A) and 50 $\mu\text{g}/\text{kg}$ (B) groups. Dramatic bile duct proliferation (arrow head) and necrosis (arrow) were found at 250 $\mu\text{g}/\text{kg}$ (C) and 1000 $\mu\text{g}/\text{kg}$ (D) groups. Hemorrhagic necrosis occurred only at 1000 $\mu\text{g}/\text{kg}$ (D, arrow). Bars represent 50 μm .

FIGURE 5.3.—Bile duct proliferation and periportal necrosis induced by 5-week treatment with AFB₁. Slight bile duct proliferation was found at 25 $\mu\text{g}/\text{kg}$ group (A) and this change is more obvious at 75 $\mu\text{g}/\text{kg}$ group (B). Periportal necrosis were also found at 10 (C) and 75 $\mu\text{g}/\text{kg}$ (D) groups (arrow), necrosis at 25 $\mu\text{g}/\text{kg}$ group was not shown here. Bars represent 50 μm .

FIGURE 5.4.—AFB₁ induced proliferation foci in liver after repeated-dose treatment. 75 $\mu\text{g}/\text{kg}$ AFB₁ treatment induced formation of proliferation focus after 4-week (A) and 5-week (B) treatment (arrow). The altered foci were characteristic of clear cytoplasm and many cells have a centrally located nucleus (clear cell foci). Bars represent 50 μm .

FIGURE 5.5.—AFB₁ induced liver GST-P⁺ cells or foci formation after single-dose treatment with AFB₁. GST-P⁺ hepatocytes were not present at 3-day (A) but 5-day (B) and 1-week (C) after single-dose treatment with 50 $\mu\text{g}/\text{kg}$ AFB₁ (arrow). 250 $\mu\text{g}/\text{kg}$ AFB₁

treatment induced GST-P⁺ cells at 3-day (D), 5-day (E) and GST-P⁺ foci at 1-week (F) after single-dose treatment (arrow). Bars represent 50 μ m.

FIGURE 5.6.—AFB₁ induced liver GST-P⁺ cells or foci formation after repeated-dose treatment with AFB₁. Three-week treatment induced liver GST-P⁺ cells at 10 μ g/kg group (A) and GST-P⁺ foci at 25 (B) and 75 μ g/kg (C) groups (arrow). Five-week treatment induced GST-P⁺ foci at 10 (D), 25 (E), and 75 μ g/kg (F) groups. Note the relative area of GST-P⁺ foci. Bars represent 50 μ m.

FIGURE 5.7.—Summary of the integrative toxicopathological findings in this study.

CHAPTER 6

AFLATOXIN B₁ MODULATES THE EXPRESSION OF PHENOTYPIC MARKERS
AND CYTOKINES BY SPLENIC LYMPHOCYTES OF MALE F344 RATS ¹

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Abstract

Aflatoxin B₁ (AFB₁) is immunotoxic to animals and a suspicious immunosuppressant in humans. In this study we investigated the effects of AFB₁ on splenic lymphocyte phenotypes and the inflammatory cytokines expression in male F344 rats. In addition, serum AFB₁-lysine adducts level and liver and spleen histology were examined to correlate with possible immune toxicities. Exposure of animals to AFB₁ (5-75 µg/kg body weight) for 1-week showed dose-dependently decreased percentages of splenic CD8⁺ T cells and CD3⁻CD8a⁺ NK cells. A general inhibition of the expressions of IL-4 and IFN-γ by CD4⁺ T cells, IL-4 and IFN-γ by CD8a⁺ cells, and TNF-α expression by NK cells was also found; however, no concurrent histological changes in liver or spleen tissues were present, suggesting acute immunosuppression without overt toxicity. Five-week exposure with AFB₁ significantly increased the percentages of CD3⁺ and CD8⁺ T cells, especially at low doses (≤ 25 µg/kg). AFB₁ treatment significantly decreased the anti-inflammatory cytokine IL-4 expression by CD4⁺ T cells and significantly increased the pro-inflammatory cytokine IFN-γ expression by CD4⁺ T cells and TNF-α expression by NK cells. These results indicated that repeated AFB₁ exposure promotes inflammatory responses by regulating cytokines expression. Consistent with this finding, liver periportal necrosis and bile duct proliferation as well as splenic lymphocyte apoptosis were found, especially at the high dose (75 µg/kg). Our data provide novel insights into the mechanisms by which AFB₁ exposure differentially modulates the cell-mediated immune responses and suggest the involvement of inflammatory response upon repeated exposure.

Keywords: aflatoxin B₁; spleen; lymphocyte; cytokine; immunotoxicity.

Introduction

Aflatoxins (AFs) are produced predominately by the fungi *Aspergillus flavus* and *Aspergillus parasiticus*. AFs are frequent contaminants of food staples, such as maize and groundnuts, and AFs exposure is a significant public health issue in Southeast Asia and Sub-Saharan Africa^{1,2}. Aflatoxin B₁ (AFB₁) is the most potent aflatoxin that is activated in the liver to its reactive metabolite, aflatoxin B₁-8,9-epoxide. This metabolite can rapidly bind with biological macromolecules such as DNA and protein to exhibit adverse effects in animals and humans³. The immune toxic effects of AFB₁ have gained great attention due to chemical-viral interactions in the developing world where such co-exposure is common⁴.

Mycotoxins, especially AFB₁ and trichothecenes, have been shown to affect the immune system in domestic and laboratory animals⁵. A considerable amount of evidence from animal studies has demonstrated that AFB₁ possesses the immunosuppressive functions^{5,6}. Cell-mediated immunity is a particular target affected by AFB₁ exposure. AFB₁ decreases splenic helper T cells and antibody response to sheep red blood cells (SRBC) in C57BL/6 mice⁷, and inhibits production of IL-2, IL-3, and IFN- γ by splenic macrophages⁸. Murine macrophages functions in phagocytosis and production of nitric oxide and TNF- α are inhibited by AFB₁ both *in vivo* and *in vitro*^{9,10}. Dietary exposure of rats to AFB₁ (0.01-1.6 ppm) produces inhibitory effects on splenic T and B lymphocytes as well as IL-1, IL-2, and IL-6 production by splenocytes¹¹. The mechanisms behind the impaired cell-immunity have been proposed to involve the inhibition synthesis at the gene or protein level^{8,12} and the installment of the cell cycle at G₂/M¹³.

The conclusion on AFB₁ induced inhibitory effects on cytokines in particular, however, deserves careful examination because of the inconsistent findings. For example, AFB₁ exposure has shown stimulating effects on the expression of pro-inflammatory cytokines such as TNF- α , IFN- γ , and IL-6 in pigs¹⁴ while the inhibition of pro-inflammatory (IL-1 β , TNF- α) and stimulating of anti-inflammatory (IL-10) cytokine mRNA expression by AFB₁ exposure have also been reported in pigs¹⁵. Even in a single feeding assay, AFB₁ containing diet has shown differential effects based on treatment duration and doses¹¹. The model (spleen vs. white blood cells, e.g.) selection and differences in exposure duration or dose might play a role in the final outcomes.

Currently available animal data are mainly focused on the overall cytokine production from a cell mixture, the relevance to the cell-specific function is not known, due to the fact that these cytokines can be produced by various cells^{16,17}. There exists a lack of data on how AFB₁ affects cell-specific cytokine expression in animal studies. Whether AFB₁-induced histological changes in liver and spleen are related to modulatory effects on splenic lymphocyte subsets and cell-specific cytokines are largely unknown. Aims of this study are to demonstrate the modulatory effects of AFB₁ on splenic lymphocyte phenotypic markers and cell-specific cytokines, and their relevance to liver and spleen histological changes in F344 rats. This study also holds the potential to provide evidence for immunosuppressive effects for future human epidemiological studies.

Materials and Methods

Materials

Aflatoxin B₁ and dimethyl sulfoxide (DMSO) were purchased from Sigma-Aldrich Inc (St. Louis, MO). Antibodies of CD28 and CD3 and fluorescence conjugated antibodies including CD3 phycoerythrin (PE) clone G4.18, CD4 allophycocyanin (APC) clone OX-35, CD8 fluorescein isothiocyanate (FITC) clone OX-8, CD8a peridinin chlorophyll protein complex (PERCP) clone OX-8, CD45RA phycoerythrin cyanin 5 (PE-Cy5) clone OX-33, IL-4 PE clone OX-81 and INF- γ FITC clone DB-1 were purchased from BD Biosciences (San Jose, CA). Fix/perm and Perm/Wash Buffer were also purchased from BD Biosciences (San Jose, CA). TNF- α FITC clone TN3-19 antibody was purchased from Abcam (Cambridge, MA). Phorbol 12-myristate 13-acetate (PMA) and fetal bovine serum (FBS) were purchased from Sigma-Aldrich (St. Louis, MO). RPMI-1640 medium were purchased from ATCC (Manassas, VA). All other solvents were of the highest grade available.

Animals

Male F344 rats (5 weeks old, 120-130g) were purchased from Harlan Laboratory (Indianapolis, IN). Animals were individually housed under a 12h/12h light/dark cycle with temperature of $22 \pm 2^\circ\text{C}$. Purified AIH 76A diet and tap water were provided *ad libitum*. Animals were acclimated for 1-week before treatment. Animal care and use and detailed animal protocol were approved by the Institutional Animal Care and Use Committee (IACUC) at the University of Georgia.

Experimental Design

Male F344 rats were randomly divided into four groups: control, 5, 25, and 75 μg AFB₁/kg body weight (BW). Rats (5 per treatment group) were orally gavaged with AFB₁ in DMSO (50 μL /100 g BW) for duration of 1-or 5-week. Animals were sacrificed at 24 h following the last dose. Upon euthanasia, spleens were aseptically removed for preparing single cell suspension, which was used for cell surface marker and intracellular cytokine staining. The parameters tested in this study were shown in Table 6.1. The selection of these markers was based on previous studies^{18,19}.

Serum Aflatoxin B₁-Lysine Adduct (AFB-Lys) Measurement

To monitor the AFB₁ exposure, the levels of serum AFB-Lys were measured from animals at each time point. The method was adopted as previously reported²⁰. Briefly, 150 μL serum was digested with pronase (pronase:total protein, 1:4, w:w) for 3 h at 37°C. The digest was further purified by a solid phase cartridge, evaporated, reconstituted, and injected into an Agilent 1200 HPLC system for quantification by fluorescence signal. Excitation and emission wavelengths were 405 and 470 nm. The limit of detection is 0.4 pg/mg albumin. Serum AFB-Lys adduct level was adjusted by serum albumin content.

Single Splenic Cell Suspension Preparation

The procedure for preparing single splenic cell suspension was modified as previously reported²¹. Approximately one half of the spleen tissues taken at necropsy were minced in 1 mL cold complete RPMI-1640 medium (RPMI-1640 containing 2 mM L-glutamine (ATCC), supplied with 10% FBS and 100 IU/mL Penicillin and 100 IU/mL streptomycin), then gently mashed through a 70 nylon mesh cell strainer and collected into 5 mL cold complete RPMI-1640 medium. The collected cells were pelleted by

centrifuge at 400g for 5 min at 4°C and supernatants were discarded. Then red blood cells were removed by incubating with lysing buffer for 5 min at room temperature (RT). The remaining cells were washed once and collected into 3 mL complete RPMI-1640 medium. Cell count was done in a hemocytometer and was adjusted to 2×10^7 /mL by volume. Cell viability was determined by Trypan Blue staining, which showed more than 90% viable cells.

Splenic Lymphocytes Phenotypic Markers Staining

Four splenic lymphocyte surface markers tested were CD3, CD4, CD8 and CD45RA. An aliquot of 1.0×10^6 splenic cells were first blocked in 100µL block buffer (CD32, 1:100) for 10 min at RT. Then cells were pelleted after centrifuging and supernatants were discarded. These cells were incubated with staining buffer (PBS including 10%FBS and 0.1%NaN₃) containing appropriately diluted fluorescence conjugated antibodies (CD3 PE: 1:200; CD4 APC: 1:100; CD8 FITC: 1:100; CD45RA PE-CY5: 1:200) for 30 min at 4°C in dark, 100 µL per tube. Single fluorescence antibody control was included. After incubation, the cell mixtures were washed once with staining buffer and then fixed in 250 µL cold 2% formaldehyde for 12 h. The fixed cell mixtures were pelleted and resuspended in 500 µL staining buffer in plastic tubes before flow cytometric analysis within 1 day.

Stimulation of Cytokine Expression

Duplicates of 1.0×10^6 cells for each sample were transferred to a 96-well flat bottom plate pre-coated with anti-CD3 antibody (1:100). The cells were pelleted by centrifuging at 250g for 4 min at 4°C and supernatants were discarded. 100 µL of stimulating buffer (complete RPMI-1640 medium supplied with anti-CD28 antibody

(1:100), brefeldin A (1:100), PMA (50 ng/mL), and ionomycin (500 ng/mL)) was added to stimulate cytokines production for 6 h in a humidified incubator (37°C, 5% CO₂).

Intracellular Cytokine Staining

To determine cell-specific intracellular cytokines expression, both surface markers and intracellular cytokines were stained. The combination of cell surface markers and cytokine markers were (1) CD4 APC + CD8a PERCP + IL-4 PE + INF- γ FITC, and (2) CD3 PE + CD8a PERCP + TNF- α FITC. The method was modified from a previous protocol²². Briefly, cells in each well were washed twice with 200 μ L cold PBS after stimulation, then were resuspended in 100 μ L staining buffer containing either CD4 APC (1:100) + CD8a PERCP (1:100) antibodies or PE CD3 (1:200) + CD8a PERCP (1:100) antibodies and incubated for 30 min at 4°C in dark. Cells were pelleted by centrifuging at 250g for 4 min at 4°C, washed once with 250 μ L staining buffer. Next, 100 μ L Fix/Perm buffer were added into each well, which was mixed quickly and thoroughly and incubated for 30 min at 4°C in dark. The cells were washed twice with 250 μ L 1 \times Perm/Wash buffer (500g, 4°C, 5 min). Then intracellular cytokines staining was followed. Conjugated anti-cytokines were prepared in 1 \times Perm/Wash buffer as follows: IL-4 PE: 1:100 and IFN- γ FITC: 1:5; TNF- α FITC: 1:100. Cells were incubated in a mixture of IL-4 PE and IFN- γ FITC or TNF- α FITC for 30 min at 4°C in dark. After intracellular staining, cells were washed twice with 1 \times Perm/Wash buffer (500g 4°C, 5 min), followed by fixation with 200 μ L cold 2% formaldehyde. Lastly, cells were washed once with staining buffer and resuspended in 500 μ L staining buffer in plastic tubes before flow cytometric analysis within 1 day.

Flow Cytometric Analysis of Splenic Lymphocyte Phenotypes and Intracellular Cytokine Expressions

A total of 20,000 cells for either surface markers or different combinations of cytokine markers were counted on a HyperCyAn flow cytometer (Beckman Coulter Inc., Fullerton, CA) for analysis of splenic lymphocyte phenotype or cytokine expression and data were analyzed by FlowJo software (Tree Star Inc., Ashland, OR). Selective histograms and dot-plot images were shown in Figure 6.1.

H&E Staining of Liver Tissues

Dissected rat liver tissues were fixed in 4% neutral formalin and then embedded in paraffin. Tissue sections (5 μ m) were cut and routinely processed for H.E. staining and examined under light microscopy. Photographs were taken on an Olympus XC30 microscope with an Olympus UC30 digital camera (Olympus America Inc., Center Valley, PA) linked to a Celsense image analysis system (Celsense Inc., Pittsburgh, PA). The apoptotic cells in splenic tissue sections were defined by the presence of heterochromatic cell fragments or apoptotic bodies, as previously described²³.

Statistical Analysis

Serum levels of AFB-Lys adduct, the percentages of surface marker and cytokine expression were expressed as mean \pm standard deviation (SD). Comparisons of these parameters between treated groups and the control group were conducted using one-way analysis of variance (ANOVA) followed by Tukey's multiple comparisons. A *p*-value of less than 0.05 was considered statistically significant.

Results

Serum Levels of AFB-Lys Adducts

As shown in Table 6.2, serum levels of AFB-Lys adduct in animals were measured to reflect the internal doses. Dose-dependent increases of serum AFB-Lys adduct levels were found after 1- or 5-week treatment. The adduct levels increased by 134.95% and 98.41% after 5-week treatment for the 5 and 25 $\mu\text{g}/\text{kg}$ groups, respectively. However, only slight change in the adduct level was found for the 75 $\mu\text{g}/\text{kg}$ group compared to that after 1-week treatment (7.32 vs. 7.18 ng/mg albumin). The typical chromatograms were shown in Figure 6.2.

Effects of 1-week Exposure of AFB₁ on the Expression of Splenic Lymphocytes Phenotypic Markers and Intracellular Cytokines

The effects of AFB₁ on the percentage of splenic lymphocyte subsets after 1-week treatment are shown in Figure 6.3. Dose-dependent decreases in the percentages of cytotoxic T cells (CD8⁺) and CD3⁻CD8a⁺ NK cells were found after 1-week treatment, and statistical significance was found for the 25 and 75 $\mu\text{g}/\text{kg}$ groups as compared to the control ($p < 0.05$). No statistically significant effects on T lymphocytes (CD3⁺), helper T lymphocytes (CD4⁺), or B lymphocytes (CD45RA⁺) were found after 1-week treatment (data not shown).

AFB₁ treatment produced a dose-related and significant reduction of IL-4 expression by CD4⁺ T cells at all dose levels after *ex vivo* stimulation ($p < 0.05$, Figure 6.4A). Dose-dependent decrease of IFN- γ expression by CD4⁺ T cells was also found with significant difference only at 75 $\mu\text{g}/\text{kg}$ group ($p < 0.05$, Figure 6.4B). A significant decrease of IL-4 expression by CD8a⁺ cells occurred at 25 and 75 $\mu\text{g}/\text{kg}$ groups ($p <$

0.05), with 39% and 58% decrease, respectively (Figure 6.4C). Decreased IFN- γ expressions by CD8a⁺ cells also appeared at 25 and 75 $\mu\text{g}/\text{kg}$ groups ($p < 0.05$, Figure 6.4D). The effects of AFB₁ on IL-4 and IFN- γ expression by CD8a⁺ cells also exhibited a dose-related change. Dose-dependent decrease in the expression of TNF- α by CD3⁻ CD8a⁺ NK cells was found with the statistically significant difference observed only at 75 $\mu\text{g}/\text{kg}$ group ($p < 0.05$, Figure 6.4E).

Effects of 5-Week exposure of AFB₁ on the Expression of Splenic Lymphocytes Phenotypic Markers and Intracellular Cytokines

The effects of AFB₁ treatment on splenic lymphocyte phenotypes after 5-week treatment are shown in Figure 6.5. Significant increases in the percentages of CD3⁺ and CD8⁺ T lymphocytes were observed at 5 and 25 $\mu\text{g}/\text{kg}$ groups ($p < 0.05$) but not at 75 $\mu\text{g}/\text{kg}$ group ($p > 0.05$). Other splenic lymphocyte subsets including CD4⁺ T cells, B cells and NK cells were not significantly affected after 5-week treatment with AFB₁ (data not shown).

Effects of AFB₁ on cytokine expression are shown in Figure 6.5. Differential effects on IL-4 and IFN- γ expressions by CD4⁺ T cells were found. Significant decreases in percentages of IL-4 expressing CD4⁺ T cells were found at all dose levels ($p < 0.05$, Figure 6.6A) while a significant increase in IFN- γ expressing CD4⁺ T cells (37.8%) occurred only at 25 $\mu\text{g}/\text{kg}$ group ($p < 0.05$, Figure 6.6B). A significant decrease of IL-4 expressing CD8a⁺ cells was also found at 25 $\mu\text{g}/\text{kg}$ group ($p < 0.05$, Figure 6.6C). Although the inhibition at 75 $\mu\text{g}/\text{kg}$ group was not statistically significant, the fluorescence intensity of IL-4 in CD8a⁺ cells was significantly decreased by 58% ($p <$

0.05, data not shown). In addition, there was also a significant elevation of TNF- α expression by CD3⁻CD8a⁺ NK cells (85.9%) at 75 μ g/kg group ($p < 0.05$, Figure 6.6D).

The pattern changes of splenic lymphocyte phenotypes and cell-specific cytokine expressions after 1-week and 5-week AFB₁ exposure are summarized in Table 6.3. IL-4 expression by CD4⁺ T cells is consistently decreased following both 1-week and 5-week exposure; IFN- γ expression by CD4⁺ T cells and TNF- α expression by NK cells experienced a temporal inversion from being decreased to elevation.

Effects of AFB₁ on Liver and Spleen Histology

One-week treatment with AFB₁ did not cause apparent histological changes at any dose group (data not shown). However, repeated treatment for 5-week induced dose-related periportal necrosis in the liver (Figure 6.7). In addition, bile duct proliferation was apparent at 25 and 75 μ g/kg groups, with much larger area occupied by proliferated bile duct at 75 μ g/kg group (Figures 6.7E and F); however, no bile duct proliferation was induced at 5 μ g/kg group (Figure 6.7B).

No apparent histological changes of the splenic microstructures were found after 1-week or 5-week treatment with AFB₁ at all dose levels (Figure 6.8) except that treatment with 75 μ g/kg AFB₁ caused an increased apoptosis of lymphocytes after 5-week (Figure 6.8D).

Discussion

Results of this study demonstrated that short-term exposure (1-week) to AFB₁ could dose-dependently decrease the percentages of splenic CD8⁺ lymphocytes and NK cells and the production of cytokines by CD4⁺, CD8⁺ lymphocytes and NK cells; prolonged exposure (5-week) to low levels of AFB₁ ($\leq 25\mu$ g/kg) had stimulatory effects

on splenic lymphocytes but not at the high dose (75 $\mu\text{g}/\text{kg}$); prolonged 5-week exposure also decreased IL-4 expression while increasing IFN- γ and TNF- α expression. Splenic B lymphocytes were not significantly affected throughout the experimental period at all dose levels. Data from this study provide evidence of the modulatory effects of AFB₁ on splenic lymphocyte phenotypes and cell-specific cytokine expression. To our knowledge, this is the first animal study to investigate the effects of AFB₁ on lymphocyte specific functions in cytokine production. Results of this study demonstrate that both splenic lymphocyte phenotypes and cell specific cytokine production can be modulated by AFB₁, and prolonged exposure may play a role in chronic inflammation through regulating cytokine expression.

Doses selected in this study were comparable to human exposure which were approximately equivalent to 0.175, 0.875, and 2.625 ppm, respectively, in contaminated food. The doses herein are relevant to human exposure especially for populations in developing countries where food contamination is a common problem. For instance, the individual daily exposures can be expected to range from 1 to more than 50 mg AFB₁ in Southern China and Kenya^{24, 25}.

Serum AFB-Lys adduct is an established biomarker for AFB₁ exposure²⁶, and it was adopted to monitor the animal dosimetry for exposure in this study. Serum AFB-Lys adduct levels exhibited different patterns: at doses $\leq 25\mu\text{g}/\text{kg}$, it increased by about 2 fold after 5-week treatment compared to that after 1-week treatment; nevertheless, there was no obvious change for the 75 $\mu\text{g}/\text{kg}$ group. This observation may indicate a plateau reached for this biomarker or a damaged metabolic balance for liver enzyme functions in study animals. The periportal necrosis and bile duct proliferation found in liver at this

dose group supports that the liver injury influenced AFB₁ metabolic activation and adduct formation.

CD4⁺ and CD8⁺ T lymphocytes are important effector cells for cell-mediated immune responses²⁷. NK cells serve as a defensive barrier to kill pathogens and foreign cells and participate in the innate and adaptive immune mechanisms²⁸. CD8⁺ T cells from spleen of normal rats almost exclusively express the alpha/beta isoform while NK cells carry only CD8 alpha chains (alpha/alpha)^{18, 29}. Therefore, splenic CD8a⁺ cells could be a mixture of both CD8⁺T cells and CD3⁻CD8a⁺NK cells, with the majority of which being CD8⁺ cell. Due to the limited commercial availability of conjugated antibodies for rat, the CD8a⁺ cells were chosen for assessing cytokine expression in this study. These lymphocytes secrete several different cytokines such as IL-4, INF- γ and TNF- α , which are important coordinators of immune and inflammatory responses³⁰. In this study, we found that acute exposure to AFB₁ for 1-week decreased the percentages of CD8⁺ lymphocytes and NK cells. The impaired cell-mediated immunity was similarly observed in other species such as mice³¹ and chickens³². In agreement with this, B lymphocytes (CD45RA⁺) were not significantly affected by AFB₁ after 1-week. Although the percentage of CD4⁺ T lymphocytes was not reduced, the expression of IL-4 and IFN- γ by CD4⁺ T lymphocytes was significantly decreased at the high dose (75 μ g/kg). On the other hand, both the percentage of CD8⁺ T lymphocytes and the intracellular expression of IL-4 and IFN- γ were significantly decreased after 1-week treatment, indicating that CD8⁺ T lymphocytes might be more susceptible to AFB₁ exposure. TNF- α expression by NK cells was also found to be inhibited after 1-week exposure with AFB₁. Apparently these immune effects were not relevant to overt toxicity in liver or spleen tissues as

evidenced by histological results. Instead, the inhibitory effects could result from inhibited synthesis at either gene or protein level or restricted cell cycle at G₂/M, which has been reported by other studies^{8,12,13}.

Dose-dependent effects on splenic lymphocyte phenotypes were found after 5-week prolonged exposure, e.g., lower doses of AFB₁ (5 and 25 µg/kg) increased percentages of CD3⁺ and CD8⁺ T lymphocytes, while the highest dose (75 µg/kg) had no apparent effect. This finding was similar to deoxynivalenol (DON) induced immune modulations that low dose DON exposure upregulates cytokines and inflammatory genes with concurrent immune stimulation whereas high dose enhances leukocyte apoptosis with concurrent immune suppression³³. Earlier studies have also demonstrated that repeated treatment with AFB₁ for 4-week could enhance protein and DNA synthesis in splenic lymphocytes at low dose (30 µg/kg body weight), but not at high doses (145 µg/kg body)¹², which is in agreement with our findings.

Contrary to the overall inhibition of cytokine production after 1-week exposure to AFB₁, significant inhibition of IL-4 expression by CD4⁺ and CD8⁺ cells and stimulation of IFN-γ expression by CD4⁺ cells and TNF-α expression by NK cells were found after 5-week exposure to AFB₁. Stimulation of INF-γ production by CD4⁺ cells could be a direct response or a feedback caused by inhibition of IL-4 given the interaction between IL-4 and INF-γ¹⁶. The stimulation of TNF-α production by NK cells might take part in chronic inflammation, which is consistent with the elevated IL-6 production as reported previously¹¹ since TNF-α can stimulate splenic production of IL-6 in rat³⁴. Given the anti-inflammatory effect of IL-4 and pro-inflammatory effects of INF-γ and TNF-α³⁵, the inhibition of IL-4 and stimulation of INF-γ and TNF-α might lead to a net result of

inflammatory responses. Data from in vitro studies also support this finding. Murine macrophages exposed to AFB₁ showed decreased secretion of anti-inflammatory IL-10 and increased pro-inflammatory IL-6³⁶ and similar effects were found in fumonisinB₁ treated pig peripheral blood mononuclear cells which demonstrated an inhibitory expression of IL-4 and a stimulation of INF- γ at both protein and mRNA levels³⁷. Considering the rather short turnover time of these cytokines³⁸, it seems unlikely that the observed pattern on cytokines are due to this natural turnover. However, further research is warranted to clarify the potential mechanisms involved and the complications associated with cytokine interactions, such as direct versus indirect effects.

In this study the dose-related induction of liver periportal necrosis and bile duct proliferation was indicative of inflammatory responses in liver and in agreement with the altered profiles of cytokines after 5-week exposure to AFB₁. Prolonged exposure also induced slightly increased apoptosis of splenic T lymphocytes at the high dose (75 $\mu\text{g}/\text{kg}$), and this alteration was consistent with the profile changes of cytokines, such as elevated TNF- α expression and INF- γ expression. Synergistic action of TNF- α and INF- γ in promoting apoptosis of pancreatic beta cells has been reported³⁹, and mode of action might be through activating JNK/SAPK and p53 pathway with the involvement of reactive oxygen species (ROS) production⁴⁰.

Findings in this study indicated that the functions of CD4⁺ and CD8⁺ cells were impaired by AFB₁, which supports the finding in humans that CD8⁺ lymphocyte function was negatively correlated with AFB₁ exposure levels⁴; however, different effects were found in regard to expression of IFN- γ and IL-4 by CD4⁺ cells. The variation might be due to different experimental design and exposure period as well as different comparison

groups (non-treated rats versus widespread low exposed humans). An *in vitro* study has shown that IL-4 acts synergistically with IL-5 to increase IgA production by LPS activated murine B cells, while IFN- γ inhibits IL-5 mediated up-regulation of IgA secretion ⁴¹. Therefore, the observed inhibition of IL-4 and stimulation of IFN- γ in this study supports the finding of decreased saliva IgA in Gambian children with high AFB₁ exposure ⁴².

To sum up findings in the current study, a flow chart was constructed as shown in Figure 6.9. Based on previous knowledge and our data, short-term exposure to AFB₁ may elicit immunosuppressive effects through inhibiting effects on gene or protein synthesis while prolonged exposure may upregulate cytokines and proinflammatory genes to enhance inflammation and apoptosis. The data provides novel insight into AFB₁ exposure induced alterations of cell-mediated immune responses. However, exact mechanisms for the complex regulation of cytokine production, interaction between liver and spleen changes, as well as for the transition from short-term to prolonged effects need further studies to clarify.

Conflict of interest statement

The authors declare no conflict of interest in this work.

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Table 6.1. Immune parameters examined.

Groups (cellular expression)	Immune parameters
T-lymphocytes	CD3 ⁺
T cell subsets	CD4 ⁺ , CD8 ⁺
B cells	CD45RA ⁺
NK cells	CD3 ⁻ CD8a ⁺
Cytokine expression	(IL-4-) expression by CD4 ⁺ or CD8a ⁺ T cells, (IFN- γ -) expression by CD4 ⁺ or CD8a ⁺ T cells, (TNF- α -) expression by CD3 ⁻ CD8a ⁺ NK cells

Table 6.2. Serum levels of AFB-Lys adduct.

Dose ($\mu\text{g}/\text{kg}$)	Serum AFB-Lys levels (ng/mg albumin)	
	1-week	5-week
0	ND	ND
5	0.22 \pm 0.01	0.52 \pm 0.02 ^c
25	1.54 \pm 0.09 ^a	3.06 \pm 0.15 ^{a,c}
75	7.18 \pm 0.28 ^{a,b}	7.32 \pm 0.28 ^{a,b}

Note. Values are mean \pm SD for 5 rats. ND, not detectable. ^a $p < 0.001$, compared to 5 $\mu\text{g}/\text{kg}$ in the same column. ^b $p < 0.001$, compared to 25 $\mu\text{g}/\text{kg}$ in the same column. ^c $p < 0.01$, compared to 1-week in the same row.

Table 6.3. Summary of the immune parameter changes.

Immune parameters (%)	1-week ($\mu\text{g}/\text{kg}$)			5-week ($\mu\text{g}/\text{kg}$)		
	5	25	75	5	25	75
CD3 ⁺ T cells	-	-	-	↑	↑	-
CD8 ⁺ T cells	-	↓	↓	↑	↑	-
CD3 ⁻ CD8a ⁺ NK cells	-	↓	↓	-	-	-
CD4 ⁺ IL-4 ⁺ T cells	↓	↓	↓	↓	↓	↓
CD4 ⁺ IFN- γ ⁺ T cells	-	-	↓	-	↑	-
CD8a ⁺ IL-4 ⁺ T cells	-	↓	↓	-	↓	-
CD8a ⁺ IFN- γ ⁺ T cells	-	↓	↓	-	-	-
CD3 ⁻ CD8a ⁺ TNF- α ⁺ NK cells	-	-	↓	-	-	↑

Note. ↑: $P < 0.05$, significant increase compared to control; ↓: $P < 0.05$, significant decrease compared to control; -: $P > 0.05$, no significant difference compared to control.

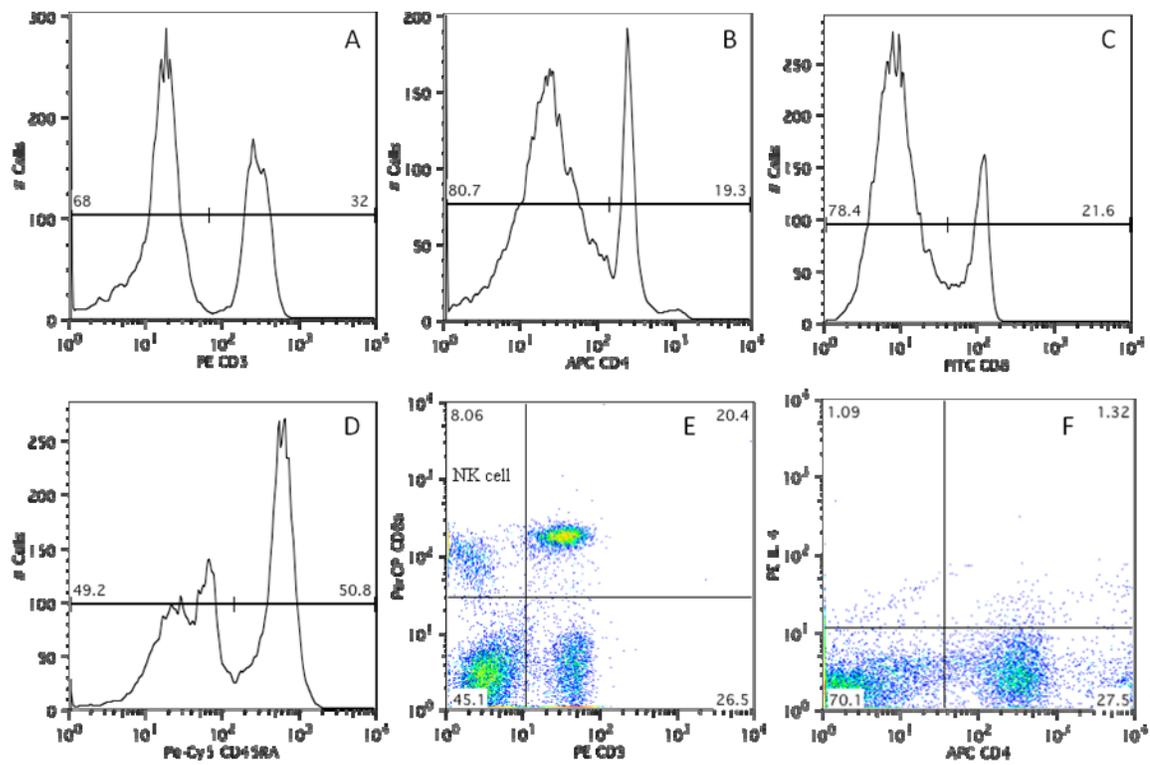


Figure 6.1. Selective typical histograms and two-parameter dot-plots.

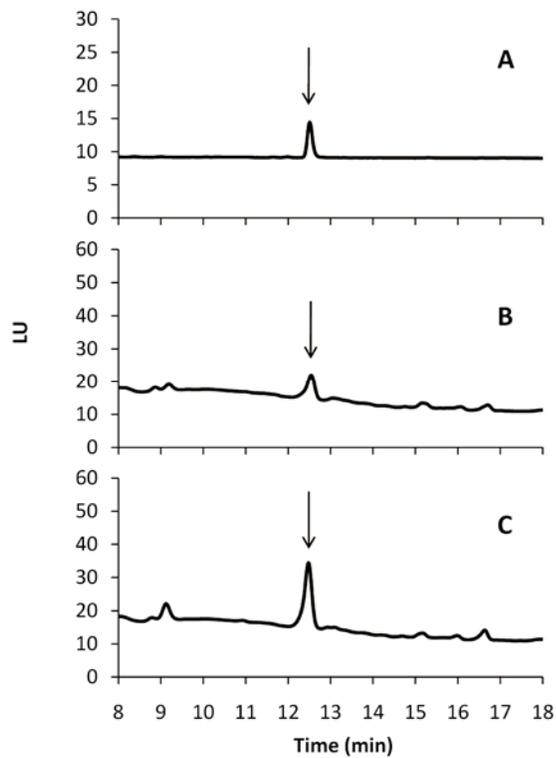


Figure 6.2. Typical chromatograms of AFB-Lys adduct.

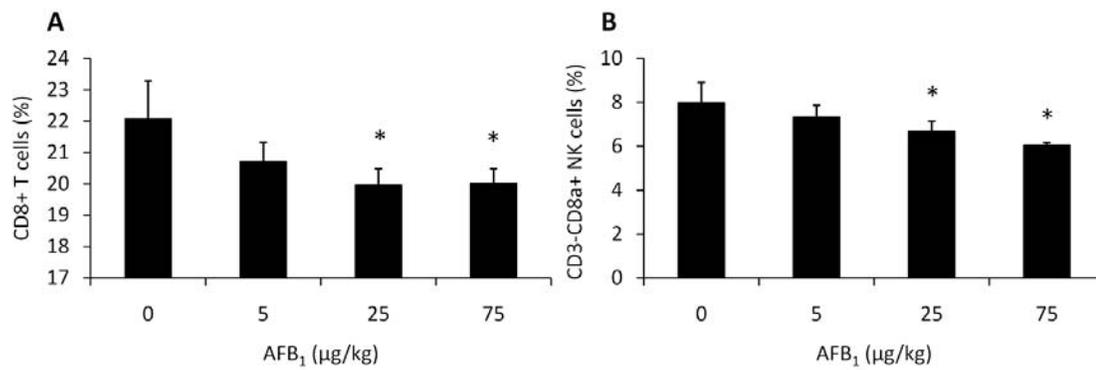


Figure 6.3. Effects of AFB₁ on splenic lymphocyte phenotypes after 1-week exposure.

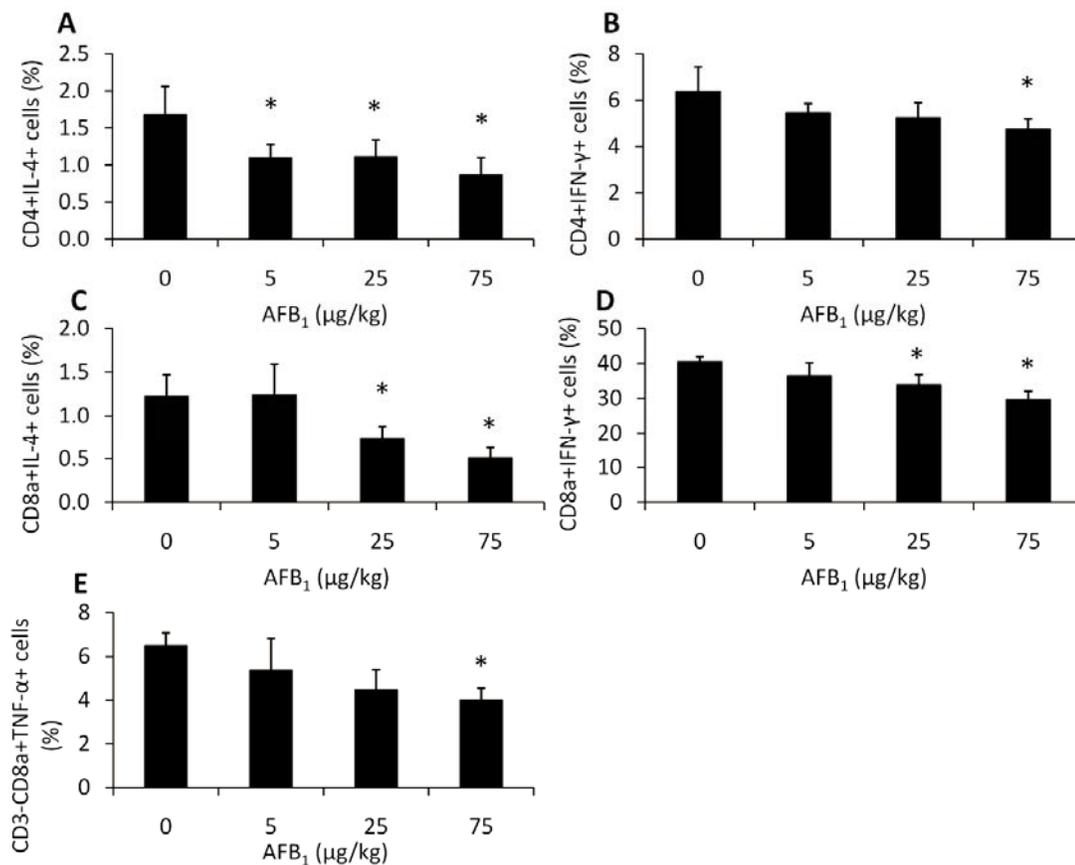


Figure 6.4. Effects of AFB₁ on cytokine expression in splenic lymphocytes after 1-week exposure.

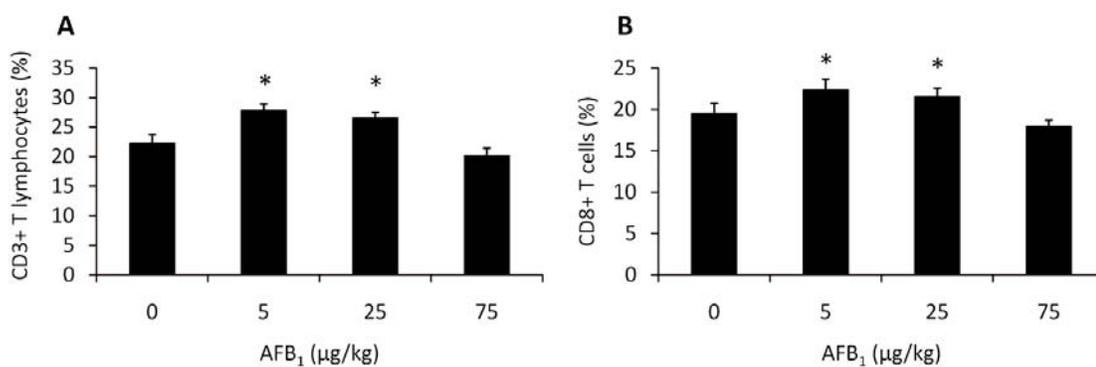


Figure 6.5. Effects of AFB₁ on splenic lymphocyte phenotypes after 5-week exposure.

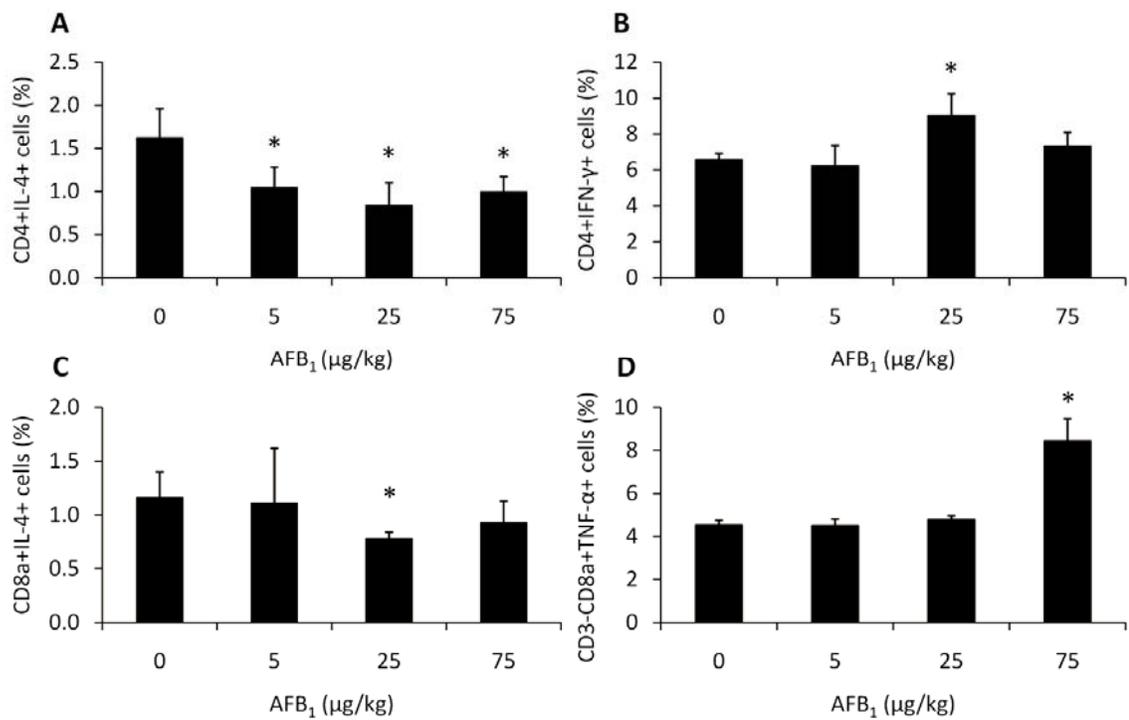


Figure 6.6. Effects of AFB₁ on cytokine expression in splenic lymphocytes after 5-week exposure.

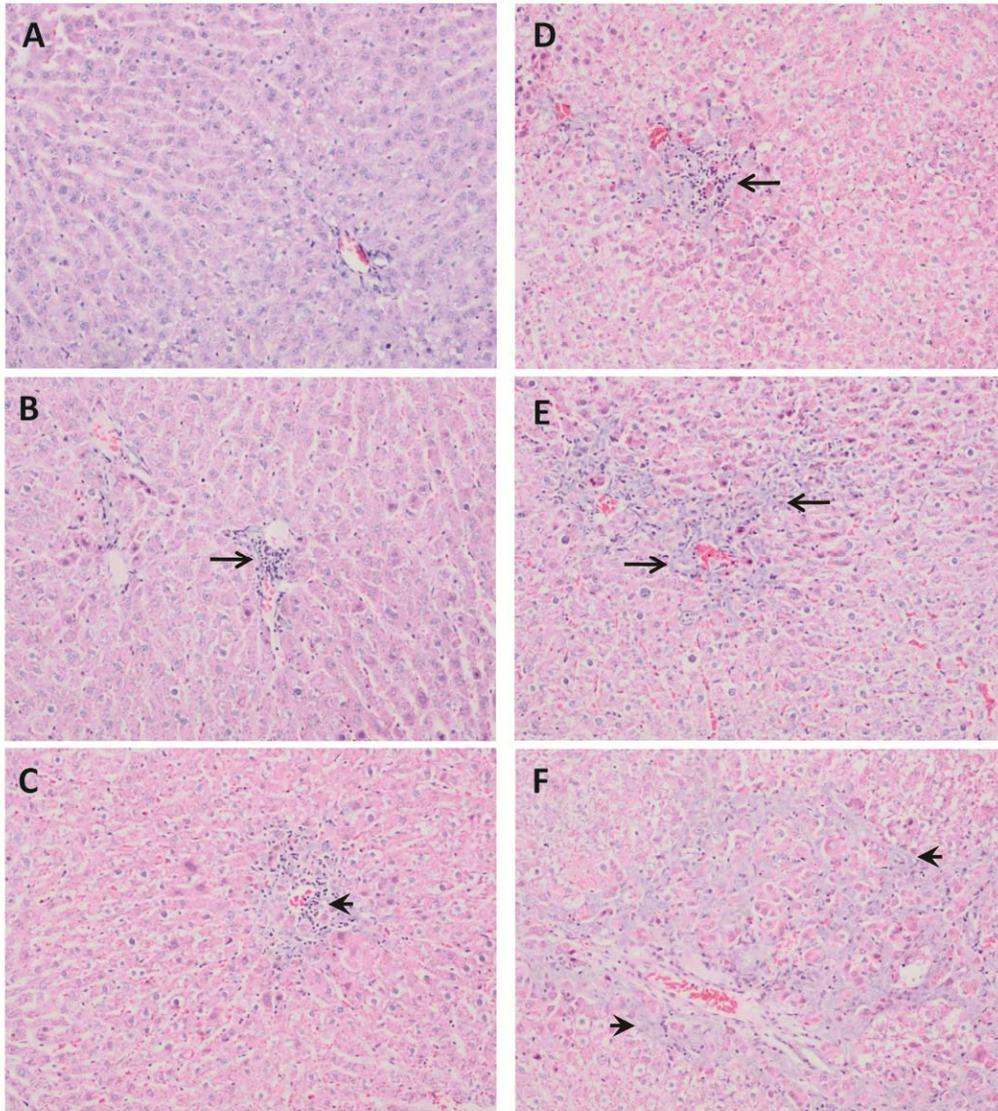


Figure 6.7. Histological effects of AFB₁ on liver after 5-week exposure.

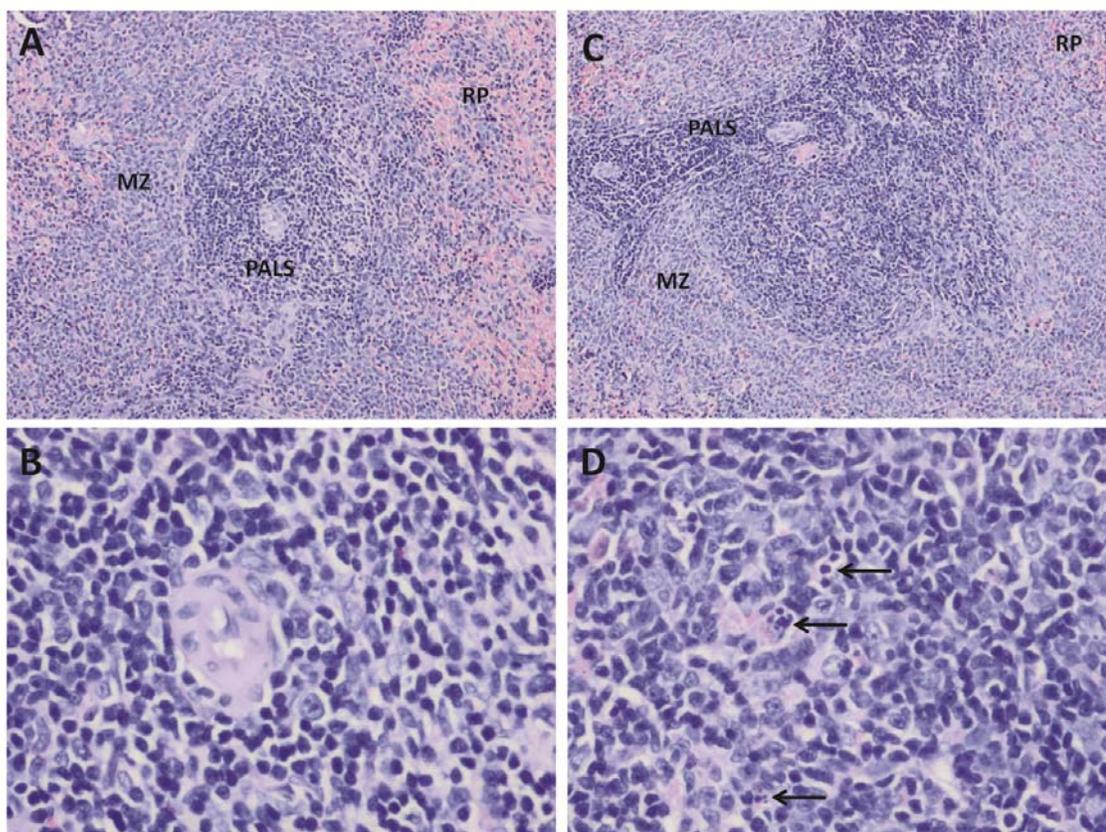


Figure 6.8. Histological effects of AFB₁ on spleen after 5-week exposure.

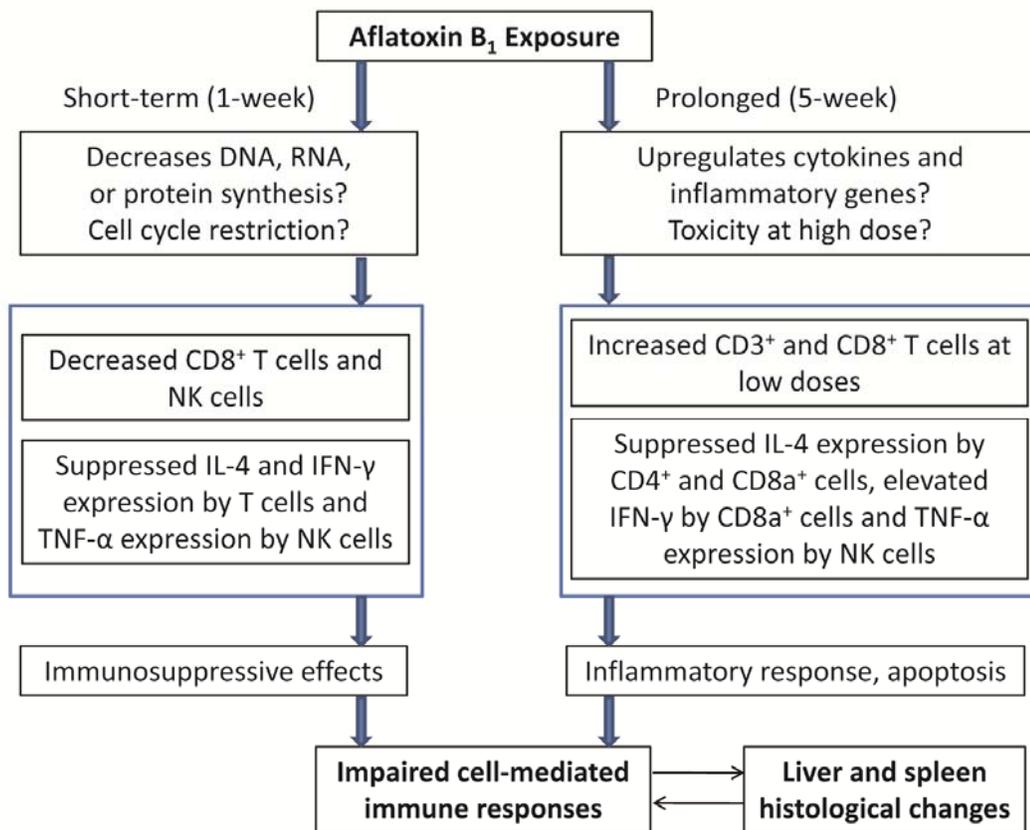


Figure 6.9. Possible mechanisms of AFB₁ immunotoxic effects in F344 rats.

Figure Legends

Figure 6.1. Selective typical histograms and two-parameter dot-plots. Histograms are PE-CD3 (A), APC-CD4 (B), FITC-CD8 (C), and Pe-Cy5-CD45RA (D). Two-parameter dot-plots show NK cells (CD3⁻CD8a⁺, E) and IL-4⁺CD4⁺ cells (F).

Figure 6.2. Typical chromatograms of AFB-Lys adduct. (A) An authentic standard, 1.0 ng/mL. (B) A diluted rat serum sample (5 fold) from 5 µg/kg AFB₁ exposure. (C) A diluted rat serum sample (10 fold) from 25 µg/kg AFB₁ exposure. Arrows indicate the peak for AFB-Lys adduct.

Figure 6.3. Effects of AFB₁ on splenic lymphocyte phenotypes after 1-week exposure. Dose-dependent decreases of the percentages of CD8⁺ T cells (A) and CD3⁻CD8a⁺ NK cells (B) were found. *Indicates $p < 0.05$ compared with control group.

Figure 6.4. Effects of AFB₁ on cytokine expression in splenic lymphocytes after 1-week exposure. Short-term exposure significantly decreases the expressions of IL-4 (A) and IFN-γ (B) by CD4⁺ T cells, IL-4 (C) and IFN-γ (D) by CD8a⁺ T cells, and the TNF-α by CD3⁻CD8a⁺ NK cells (E). *Indicates $p < 0.05$ compared with control group.

Figure 6.5. Effects of AFB₁ on splenic lymphocyte phenotypes after 5-week exposure. Long-term exposure increased the percentages of CD3⁺ T cells (A) and CD8⁺ T cells (B), especially at low dose levels. *Indicates $p < 0.05$ compared with control group.

Figure 6.6. Effects of AFB₁ on cytokine expression in splenic lymphocytes after 5-week exposure. The expression of IL-4 by CD4⁺ T cells was inhibited at all doses (A) while IFN-γ expression was elevated at 25 µg/kg group (B). The IL-4 expression by CD8a⁺ T cells was inhibited at 25 µg/kg group (C) and TNF-α expression by CD3⁻CD8a⁺ NK cells was increased at 75 µg/kg group (D). *Indicates $p < 0.05$ compared with control group.

Figure 6.7. Histological effects of AFB₁ on liver after 5-week exposure. H&E-stained liver section showed normal microstructure in control group (A). Progressed periportal necrosis was found at 5 (B), 25 (C) and 75 (D) $\mu\text{g}/\text{kg}$ groups (arrow). Bile duct proliferation occurred at 25 (E) and 75 (F) $\mu\text{g}/\text{kg}$ groups (arrowhead). Note the larger area with bile duct proliferation at 75 $\mu\text{g}/\text{kg}$ group (F) compared to 25 $\mu\text{g}/\text{kg}$ group (E). Magnification, $\times 100$.

Figure 6.8. Histological effects of AFB₁ on spleen after 5-week exposure. (A) and (C) are spleen tissues from control group at low ($\times 100$) and high ($\times 400$) magnifications. Normal spleen tissue is composed of two major parts: white pulp and red pulp (RP). The white pulp has three characteristic components: periarteriolar lymphoid sheath (PALS), follicle and marginal zone (MZ). (B) and (D) are spleen tissues from 75 $\mu\text{g}/\text{kg}$ group at low ($\times 100$) and high ($\times 400$) magnifications. Increased apoptosis was found in the PALS region at 75 $\mu\text{g}/\text{kg}$ AFB₁ treated spleen tissue, the arrows indicate small heterochromatic cell fragments, indicative of apoptotic bodies.

Figure 6.9. Possible mechanisms of AFB₁ immunotoxic effects in F344 rats.

CHAPTER 7
GLOBAL EXPOSURE OF AFLATOXIN B₁ IN DIFFERENT HUMAN
POPULATIONS ¹

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Abstract

Aflatoxin exposure is a significant health issue worldwide. Assessing the current exposure status in different human populations is beneficial for effective chemoprevention strategies. OBJECTIVES: This study aims to reveal the current human exposure status to aflatoxins using the molecular biomarker, serum aflatoxin B₁-lysine (AFB-Lys) adducts. In this study we assessed the global human exposure to aflatoxins using a reliable exposure biomarker, AFB-Lys adduct in serum. We recruited serum samples from populations in West African countries (Ghana and Burkina Faso), an East African country (Uganda), Southeast Asian countries (China and Malaysia), a Caribbean country (Haiti) and the United States. The biomarker levels were determined by a non-antibody based HPLC method with fluorescence detection. Varied levels of AFB-Lys were found globally and the highest exposure was found in Burkina Faso (29.89 ± 74.16 pg/mg albumin) and the lowest in the United States (0.89 ± 2.08 pg/mg albumin). Consistent with this finding, the detection rate is highest in Burkina Faso (100%) and lowest in the United States (17.2%). The rank of exposure based on the biomarker levels from high to low is: West Africa countries > Southeast Countries > Uganda > Caribbean country > the United States. Currently AFB₁ exposure remains prevalent in parts of West Africa and Southeast Asia, which warrants urgent and effective interventions.

Key words: aflatoxin, exposure, aflatoxin B₁-lysine adduct, biomarker.

Introduction

Mycotoxins are estimated to contaminate 25% of cereal crops globally¹. Among these mycotoxins, of biological significance are aflatoxins, fumonisins, deoxynivalenone, T-2, zearalenone and ochratoxins. Aflatoxins are frequent food contaminants produced mainly by *Aspergillus flavus* and *A. parasiticus* as secondary metabolites. Among the naturally occurring aflatoxins (B₁, B₂, G₁, G₂), aflatoxin B₁ (AFB₁) is most common and of predominant biological significance. The consumption of AFB₁ contaminated foods or dairy products can cause acute poisoning (aflatoxicosis) and chronic liver cancer in humans². A quantitative risk assessment has estimated that approximately 4.8-25.2% of global annual HCC cases are attributed to aflatoxin exposure alone³. The ecological role of aflatoxins has also been associated with many other adverse health effects in humans, including impaired childhood growth⁴ and immune functions⁵. In the developing world where aflatoxin exposure prevails, it was estimated that more than 40% of all disease burden is associated with aflatoxins⁶.

Traditionally, questionnaires and food surveys are common approaches to assess human exposure to food contaminants. Through the years the aflatoxin contamination of food has been frequently reported especially in sub-Saharan Africa and southeast Asia⁷. These data provide important information about the environmental aflatoxin exposure. However, the heterogeneous distribution of aflatoxins in food limits the application of food survey based intake estimation as an effective exposure measurement. Besides, the individual exposure or risk cannot be adequately estimated by multiplying the contamination level with the amount of intake, considering the inter-individual genetic variation in the metabolism and detoxification enzymes that were shown to be associated

with different risks for HCC ⁸. Instead, chemical-specific biomarkers from biological media provide an alternative measurement of individual exposure. The application of molecular biomarkers related to AFB₁ greatly facilitated the exposure assessment as well as efficacy evaluation of several chemoprevention strategies ⁹⁻¹¹.

Accumulated knowledge from toxicological and biochemical studies has revealed several valuable biomarkers related to AFB₁ exposure or risk. In the human liver, cytochrome P450 enzymes (1A2 and 3A4) are responsible for the oxidation and activation of AFB₁ to its reactive metabolite, AFB₁-8, 9-epoxide ¹². This metabolite can readily bind to macromolecules such as protein or DNA to form albumin or DNA adducts. AFB₁-albumin adducts are produced initially in liver and are transferred to the blood stream which release AFB₁-lysine (AFB-Lys) adducts under enzymatic digestion. AFB-Lys is shown to be a more specific biomarker of AFB₁ exposure compared to the albumin adducts as lysine in albumin is the only group of AFB₁ binding to a significant level ¹³. The measure of this biomarker could reflect human chronic exposure to AFB₁ for up to several months ^{14, 15}. This adduct has been used to verify the etiological role of AFB₁ exposure during the outbreak of aflatoxicosis in Kenya in 2004 ¹⁶ and for indicating human exposure to AFB₁ in several cross-sectional ¹⁷⁻¹⁹ and case-control studies ^{16, 20}. Besides, urinary metabolite AFM₁, phase II metabolite AFB₁-mecapturic acid (AFB-NAC), and the DNA adduct AFB₁-N⁷-guanine represent short-term biomarkers that correlate with AFB₁ exposure and have been used to indicate the efficacy of different intervention approaches ²¹. These short-term biomarkers were also thought to be more appropriate in evaluating the adverse immune effects of aflatoxins given the short recovery process of the immune function changes ⁶.

Various methodologies have been developed as measurement tools for serum AFB-Lys adducts. Enzyme-linked immunosorbent assay (ELISA), immunoaffinity column based high performance liquid chromatography with fluorescence detection (HPLC-f), radioimmunoassay (RIA) and isotope dilution mass spectrometry (IDMS) method form the available techniques with an addition of the solid phase extraction based HPLC-f method ²²⁻²⁶. Different methods tend to give different results, e.g. the ELISA method appears to measure some unspecific albumin adducts and produce higher estimation compared to the IDMS method ²⁷. The antibody based methods demand a high sensitivity of the antibody used and the results are prone to operating environment conditions. These different methodologies used to achieve the human exposure data restrict the direct comparison of the exposure data generated from different populations.

A global network for monitoring AFB₁ exposure based on the same method seems promising especially for comparison purpose. We have developed a non-antibody based method that has been validated in animal and humans studies ²⁶. Based on the method, we have measured the serum AFB-Lys adduct levels in different populations of the world with the aim of revealing the current AFB₁ exposure status, as well as providing evidence for future intervention strategies.

Materials and Methods

Study Populations

The study populations were recruited from different parts of the world, including West African countries (Ghana and Burkina Faso), an East African Country (Uganda), Southeast Asian countries (China and Malaysia), a Caribbean country (Haiti) and the United States. The distribution of these study populations were indicated in Figure 7.1. Burkina Faso study participants were recruited from three villages from 2001 to 2002. Ghana study populations were pregnant women recruited from two hospitals in Kumasi in 2006. Uganda study participants were from a cohort study conducted in the Medical Research Council and these samples were collected in 2011. Guangxi HCC cases and controls were recruited from Guangxi province, China, from 2004 to 2005. Malaysia study populations were healthy adults recruited from Penang, Malaysia in 2008. San Antonio participants were enrolled from three zip codes (with high liver cancer incidence) within the San Antonio metropolitan area of Bexar Co. from 2007 to 2008; Lubbock participants were local healthy volunteers that were recruited in 2005.

Serum Sample Processing

Thawed human serum samples were measured for albumin and total protein concentrations using procedures as previously described²³. A portion of each serum sample (150 μ L) was digested by pronase (pronase:total protein, 1:4, w:w) at 37°C for 3 h to release AFB-Lys adducts. AFB-Lys in digests were further extracted and purified by passing through a Waters MAX™ SPE cartridge, which was preprimed with methanol and equilibrated with water. The loaded cartridge was sequentially washed with water, 70% methanol, and 100% methanol containing 1% ammonium hydroxide at a flow rate

of 1 mL/min. Purified AFB-Lys was eluted with 2% formic acid in methanol. The eluate was vacuum-dried with a Labconco Centrivap concentrator (Kansas City, MO) and reconstituted for HPLC-fluorescence detection. The average recovery rate was 80%.

HPLC Conditions for Serum AFB-Lys Adduct Analysis

The analysis was conducted using an Agilent 1200 HPLC-fluorescence system (Santa Clara, CA). The mobile phases consisted of buffer A (20 mM $\text{NH}_4\text{H}_2\text{PO}_4$, pH 7.2) and buffer B (100% Methanol). The Zorbax Eclipse XDB-C18 reverse phase column (5 micron, 4.6 x 250 mm) equipped with a guard column was used. Column temperature was maintained at 25°C during analysis, and a volume of 100 μL was injected at a flow rate of 1 mL/min. A gradient was generated to separate the AFB-Lys adduct within 25 min of injection. The adduct was detected by fluorescence at maximum excitation and emission wavelengths of 405 nm and 470 nm, respectively. Calibration curves of authentic standard were generated weekly, and the standard AFB-Lys was eluted at approximately 12.3 min. Quality assurance and quality control procedures were taken during analyses, which included simultaneous analysis of one authentic standard and a quality control sample daily. The limit of detection was 0.4 pg/mg albumin. The serum AFB-Lys levels were adjusted by albumin content.

Statistical Analysis

AFB-Lys adducts were calculated for the mean \pm SD, median, range, and the detection rate. Box-plot charts were made using the SigmaPlot 10.0 software from Systat Software Inc. (San Jose, CA). Two-sample comparisons were tested with Wilcoxon-Mann-Whitney test using the STAT 12 software from StataCorp LP (College station, TX). A *p*-value less than 0.05 was considered statistically significant.

Results

Demographic Characteristics of Study Populations

Demographic characteristics of different study populations are shown in Table 7.1. The study participants had a mean age ranging from 27 to 55. Because of the different focuses of each study design, the gender distribution varied largely. In the Guangxi case-control study, the majority of participants were male while the Ghana study enrolled exclusively young pregnant women. For the San Antonio study, fewer men were recruited than women because less male participants were available due to work commitments. The gender distribution in Uganda, Burkina Faso and Lubbock was similar and approximately equal numbers of males and females were found.

Serum AFB-Lys Adduct Levels in Study Populations

The mean \pm SD, median, range and detection rate of serum AFB-Lys adduct levels are summarized in Table 7.2. The levels of AFB-Lys varied in different study populations. West African (Burkina Faso and Ghana) and Southeast Asian countries (China and Malaysia) possessed the highest exposure, followed by the Middle African country (Uganda) and Haiti. The United States had the lowest exposure. The detection rate also supports this ranking, i.e. serum AFB-Lys adducts were most frequently detected in West African populations and least detected in American populations. Ghanaian pregnant women were found to have high levels of this adduct in their sera.

The box-plot chart of the distribution of the serum AFB-Lys adducts in different study populations is shown in Figure 7.2. It is obvious that high levels of AFB-Lys adducts were associated with more extreme values. The individual variations were most

prominent in West African countries and were least prominent in the United States populations.

Distribution Frequency of Serum AFB-Lys Adducts in Different Study Populations

The distribution frequencies of serum AFB-Lys adducts are shown in Figures 7.3a-g. Burkina Faso and Malaysia study participants had the most frequent exposure level at 5-10 pg/mg albumin. In addition, the highest exposure level (≥ 40 pg/mg albumin) was most frequently found in Burkina Faso and Ghana. Study participants from Guangxi, Ghana, Uganda and the United States had exposure levels mainly within 1-5 pg/mg albumin.

Gender Distribution of Serum AFB-Lys Adducts Levels

The level of AFB-Lys adducts varied between genders among different study populations (Table 7.3). The exposure between male and female study participants from the United States (San Antonio and Lubbock) and Uganda was comparable ($P > 0.05$). On the other hand, male participants in Guangxi, China and Burkina Faso seemed to have higher exposure compared to their corresponding female participants, although no significant differences were found ($P > 0.05$). However, the overall gender difference in the adduct levels from the pooled study participants of all study populations was statistically significant (Figure 7.4, $P < 0.0001$), with the male participants possessing higher level of the AFB-Lys adducts than female participants. The (mean \pm SD) of AFB-Lys levels were (18.62 \pm 56.72) pg/mg albumin for males and (10.62 \pm 19.17) pg/mg albumin for females. (Malaysia and Haiti data were not included into this analysis.)

Discussion

This study represents a large scope, collaborative effort to assess the current AFB₁ exposures in global populations using the dosimetry of molecular biomarker, i.e. serum AFB-Lys adduct. One highlight of this study is that all the analyses were conducted in the same laboratory which greatly facilitates the comparison of data from different populations. We demonstrate that the exposure level to AFB₁ varied greatly in different geographical areas and that a large inter-individual variance was present even in the same study population. The highest exposed populations were found in West Africa (Burkina Faso and Ghana) and Southeast Asia (Guangxi and Malaysia), followed by East Africa (Uganda) and Latin America (Haiti), and the least exposed population were found in the United States. The male participants had a mean AFB-Lys adduct level approximately 1.75-fold higher than that in female participants, as estimated from the pooled populations. The high level and the high detection rate of serum AFB-Lys adduct in West Africa and Southeast Asia strengthens the need for urgent intervention strategies especially considering the HCC risks associated with AFB₁ exposure.

The recruited study populations are mainly located between 40°N and 40°S of the equator, as described by Williams et al.⁶. The climate of this zone is featured by hot and dry weather (the dry season in Africa) and warm and humid (Southeast Asia and the rain season in Africa), which favor the growth of the toxicogenic fungi responsible for AFB₁ production either before harvest or during storage²⁸. The increasingly warming and erratic climate adds to the uncertainty of how climate change will affect the growth of aflatoxin-producing fungi as well as the insect damage which may contribute to aflatoxin contamination²⁹. The adverse health effects in humans in the high-risk areas

drive the efforts to formulate many epidemiological studies to assess the AFB₁ exposure status or to explore the causative effect of AFB₁ exposure in human diseases, such as HIV infection in Uganda (our collaborative project).

Evidence from human and animal studies has established AFB₁ exposure as a risk factor for human HCC development. In this study, the highest AFB-Lys levels were found in populations from Africa (West and Middle) and Southeast Asia (China and Malaysia). Distribution frequency analysis showed that the highest exposed populations also had larger proportions of people with higher concentrations of AFB-Lys adducts. Consistently, the highest global HCC incidences have been reported in these areas³⁰. The endemic of several viral infections (HBV and HCV) in this zone makes these geographical areas more susceptible to human HCC, due to the synergistic effects of AFB₁ exposure and chronic HBV/HCV infection in increasing HCC risks³¹.

The mechanisms of AFB₁ induced hepatocarcinogenesis have been increasingly understood. The active metabolite of AFB₁ can bind to DNA to form DNA adducts¹⁵ and preferentially induce the transversion of G to T at codon 249 of the *p53* tumor suppressor gene in human hepatocytes³². In addition, it was demonstrated that AFB₁ produces mutations at codon 12 of the *c-ras* oncogenes using DNA extracted from AFB₁-induced transformed cell lines or primary liver tumors³³. Besides the genotoxic effects, AFB₁ exposure has been shown to be associated with a global hypomethylation in HCC³⁴, suggesting a role of AFB₁ in inducing the epigenetic changes. Therefore, both genetic and epigenetic mechanisms may be involved during the development of HCC.

The issue of food contamination by AFB₁ in the above-mentioned high-risk areas has been frequently reported. The recent food surveys have reported that the estimated

level of AFB₁ in contaminated corn samples ranged 0.20-888.30 µg/kg in China (n = 279)³⁵ and 0.54-15.33 µg/kg (n = 95) in Malaysia³⁶. The ranges of AFB₁ in contaminated corn were 3.4-636 µg/kg (n = 26) in Burkina Faso³⁷ and 0.4-490.6 µg/kg (n = 40) in Ghana³⁸, respectively. These food survey based estimations of AFB₁ exposure provide valuable information about the potential external exposure and on the estimation of daily intake. These reported high food contamination to AFB₁ strongly support our finding of high AFB-Lys adduct levels in populations from these countries. However, these estimations fail to take into account the biological and social factors important for AFB₁ metabolism, and thus may be inadequate for monitoring human risks.

Over the years, our laboratory has aimed to develop and validate molecular biomarkers associated with mycotoxin exposure and toxicity particularly for aflatoxin and fumonisin. The application of these biomarkers in epidemiological studies forms our ambitious goal of establishing a network for monitoring aflatoxin exposure. One highlight of this study is that the measure of AFB-Lys adducts was based on the same method which facilitates direct comparison. For the first time we reported the human AFB₁ exposure data in Burkina Faso, Uganda and Haiti. These preliminary data gave us an idea of the current human exposure status. Amongst the populations studied, people in Burkina Faso were most severely exposed to AFB₁, with the median serum AFB-Lys of 13.8 pg/mg albumin. Although relatively high levels of food contamination by AFB₁ have been found in Burkina Faso, Uganda, and Haiti, serum AFB-Lys levels varied largely among individual study participants in these countries as found in this study. The types of staple crops seem to play a role in the human exposure condition, e.g., the study participants in Uganda mainly eat plantains (a less frequent crop for AFB₁ contamination)

and corn and showed lower exposure compared to study participants in Burkina Faso whose staple crops are corn and peanuts, two known food media prone to AFB₁ contamination. In Haiti, AFB₁ contamination in market peanuts experienced a temporal reduction from 412.5 ± 32.1 ppb in September 2006 to 125 ± 7.1 ppb in November 2006 and further to a level of 0.20 ± 0.10 ppb in January 2007³⁹. The relatively low AFB-Lys level in Haitians may reflect the beneficial effect of these interventions to reduce AFB₁ contamination, including visual-tactile sort of the shell, flotation of select kernels, immediate roasting and grinding, and monitoring³⁹. Certainly other factors might be involved, such as the season for blood drawing, the differences in gender distribution as well as in life styles such as smoking and drinking status.

Serum AFB-Lys adduct level indicates the individual exposure and reflects the outcome of genetic polymorphisms as well as dietary modulations. It is not surprising to find the large inter-individual variation of AFB-Lys levels even in the same population. Human genetic polymorphisms in AFB₁ metabolizing enzymes (CYP450 3A5, glutathione S transferase (GST) and epoxide hydrolase) are of biological significance in the carcinogenicity of AFB₁^{40, 41}. Other factors such as drugs and dietary components may also modify the formation of this adduct by induction of enzymes or simply binding with the toxin to reduce its bioavailability⁴². Ethnic factor seems to affect the AFB-Lys adduct levels, for example, Chinese and Indians are more likely to have higher levels of AFB-Lys than Malay. This rank was noted to be contrary to the nut consumption of these ethnic groups: Indians eat more nuts, followed by Malay and Chinese. As nuts are susceptible to high level of aflatoxins in Malaysia, the contradictory finding indicated a role of genetic variance in producing AFB-Lys adducts¹⁹. Local ethnic groups in Gambia

and Ghana also showed different levels of exposure to AFB₁, with the major reasons being the differences in diet staples and food processing^{43, 44}. It seems that both genetic variance and dietary factors play an important role in determining the ethnic difference in AFB₁ exposure, or even the interaction of both. Less information on the AFB-Lys level is available in the U.S. populations, which restrict an estimation of possible ethnic difference.

There was no significant difference in serum AFB-Lys levels between males and females from each individual study population. However, when pooled, the overall AFB-Lys level is higher in males than females, and a significant difference (1.75 fold) was found ($P < 0.0001$). There is no simple answer to this question with regards to the metabolic enzymes. For example, the *GSTM1-nonnull* and *GSTT1-null* genotypes were found to be associated with a higher AFB₁-albumin adduct level in males than females in a Taiwanese population⁴⁵. In addition, the current evidence suggests that CYP1A2 activity is higher in males than females and CYP3A activity seems to be higher in females⁴⁶. Taken together, both the activities and the genetic polymorphism of metabolic enzymes need to be considered for explaining the gender difference in the AFB-Lys levels. Therefore, the current finding needs to be verified in more specific designs in human studies.

The current assessment of AFB₁ exposure was mainly based on cross-sectional studies. The feature of the study design limits the causative effect analysis of AFB₁ exposure in the etiology of human diseases. Cohort studies that use serum AFB-Lys adducts as molecular dosimetry of exposure are promising for delineating the ecological contribution of chronic AFB₁ exposure in human disease burdens, particularly in the

developing world. However, very limited information is available in this regard. Instead, our preliminary data could be used as the first step toward establishing a network for monitoring global AFB₁ exposure. This dynamic network, once established, could be used for exposure assessment, etiological studies in human diseases such as childhood development and efficacy evaluation of novel intervention strategies.

Our collective efforts show that AFB₁ exposure in West Africa and Southeast Asia remains quite common at present: more than 90% of study populations have various levels of AFB-Lys adducts. The fact that only a small fraction of AFB₁ ingested is converted to AFB-Lys adducts¹³ means that the real world exposure could be underestimated. It is estimated that approximately 4.5 billion people in the developing world are at risk for chronic exposure to aflatoxins⁶. Therefore, an urgent need for effective intervention strategies is warranted. The intervention strategies for reducing aflatoxins exposure or risk have been the concerns of several reviews^{6, 21, 42}. These strategies could be undertaken at different stages, including pre-harvest (reduce toxin production, and bio-control contamination), during food processing and storage (improve environmental setting, selection, and education), and after ingestion (reduce active metabolite formation and bioavailability). Vaccination of infants against HBV and HCV to remove the interaction with AFB₁ exposure is of priority to reduce human HCC risks in the long run³¹. The evaluation of different intervention strategies has been defined as one of the four gaps: quantification of health effects and disease burden of aflatoxin exposure; evaluation of current intervention strategies and disseminate the result; disease surveillance, food monitoring, laboratory and public health responses; and development of response protocols addressing aflatoxicosis⁴⁷.

In conclusion, this study demonstrates the current aflatoxin exposure status in different populations using serum AFB-Lys adduct as a molecular biomarker. The data presented in this study show that populations in West Africa and Southeast Asia represent the highest risk population for AFB₁ exposure at present, and highlight the task as well as the urgent need for effective interventions in these high-risk areas.

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Table 7.1. Demographic characteristics of study populations.

Factor	San Antonio	Lubbock	Guangxi Cases	Guangxi Controls	Malaysia	Ghana	Burkina Faso	Uganda
Sample size	151	70	68	136	170	755	474	127
Age (years)(mean±SD)	48 ± 15	31 ± 7	43 ± 10	43 ± 7	46 ± 17	27 ± 6	43 ± 10	55 ± 16
Gender [n(%)]								
male	43 (28.5)	32(45.7)	62 (91.2)	122 (89.7)	53 (31.2)	0 (0)	258 (54.4)	72 (56.6)
female	108 (71.5)	38(54.3)	6 (8.8)	14 (10.3)	117 (68.8)	755 (100)	216 (45.6)	55 (43.3)

Table 7.2. Global serum AFB-Lys levels in different populations (pg/mg albumin).

Parameters	San Antonio	Lubbock	Guangxi Cases	Guangxi Controls	Malaysia	Ghana	Burkina Faso	Uganda	Haiti
Numbers	151	54	68	136	175	755	474	127	193
Mean ± SD	4.44 ± 3.19	0.59 ± 0.33	10.60 ± 28.21	7.34 ± 16.56	6.31 ± 3.54	10.86 ± 19.01	26.73 ± 64.21	6.22 ± 24.44	6.12 ± 14.41
Median	3.44	0.63	4.97	3.69	6.08	5.01	13.8	2.21	2.41
Range	1.64 - 16.01	0.52-0.91	0.54 - 227.13	0.40 - 175.83	0.81 - 13.67	0.44 - 286.73	0.56 - 918.72	0.41 - 253.11	0.41 - 130.39
Detection Rate (%)	17.2	9.3	91.2	97.1	98	90.7	100	95.3	67.4

Table 7.3. Gender distribution of serum AFB-Lys levels in different populations (pg/mg albumin).

Gender	San Antonio	Lubbock	Guangxi Cases	Guangxi Controls	Ghana	Burkina Faso	Uganda
Male	4.15±2.76	0.69±0.20	11.30±29.46	6.30±8.35	NA	28.71±74.70	3.07±2.18
Female	4.63±3.51	0.73±0.21	3.45±2.59	4.19±3.69	10.86±19.01	24.37±48.88	4.02±5.56
<i>p</i> -value	0.5444	1	0.1247	0.6039		0.9952	0.7694

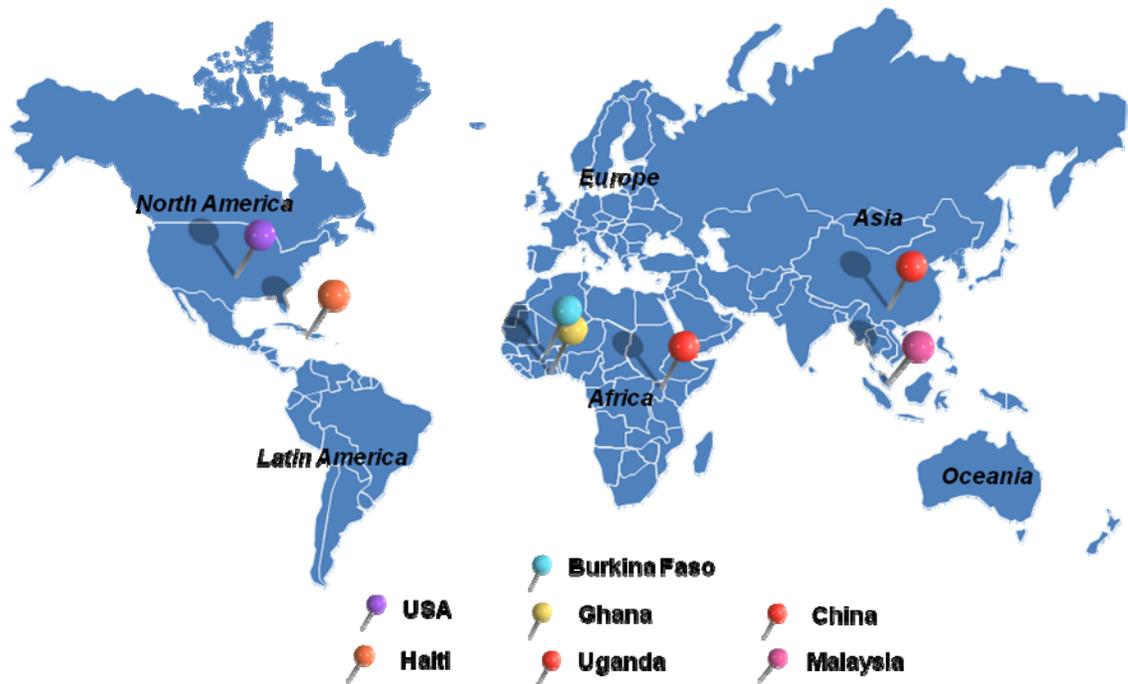


Figure 7.1. Geographic distribution of different study populations in this study.

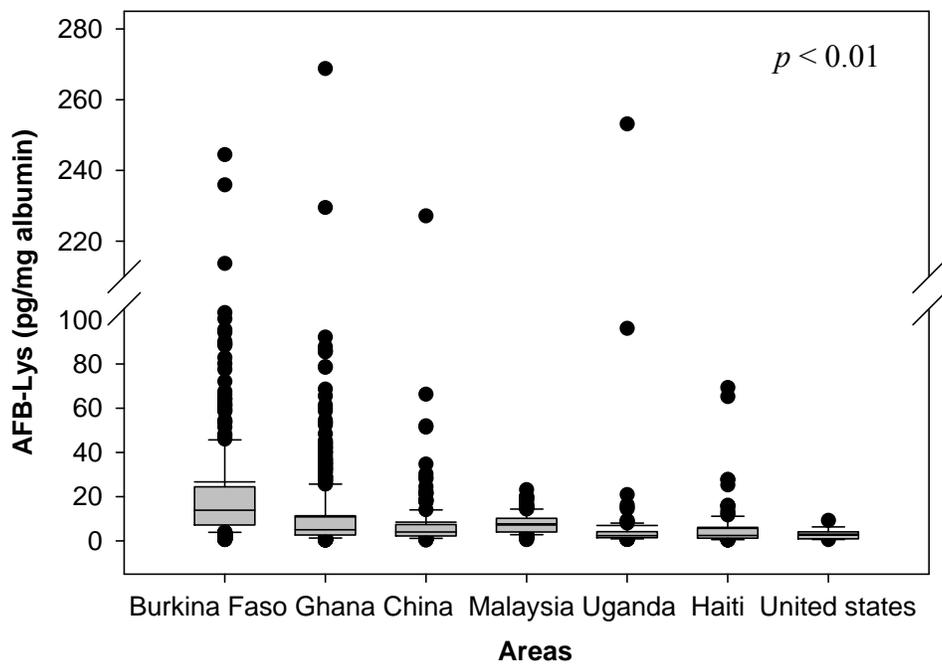


Figure 7.2. Box-plot of serum AFB-Lys levels in different study populations.

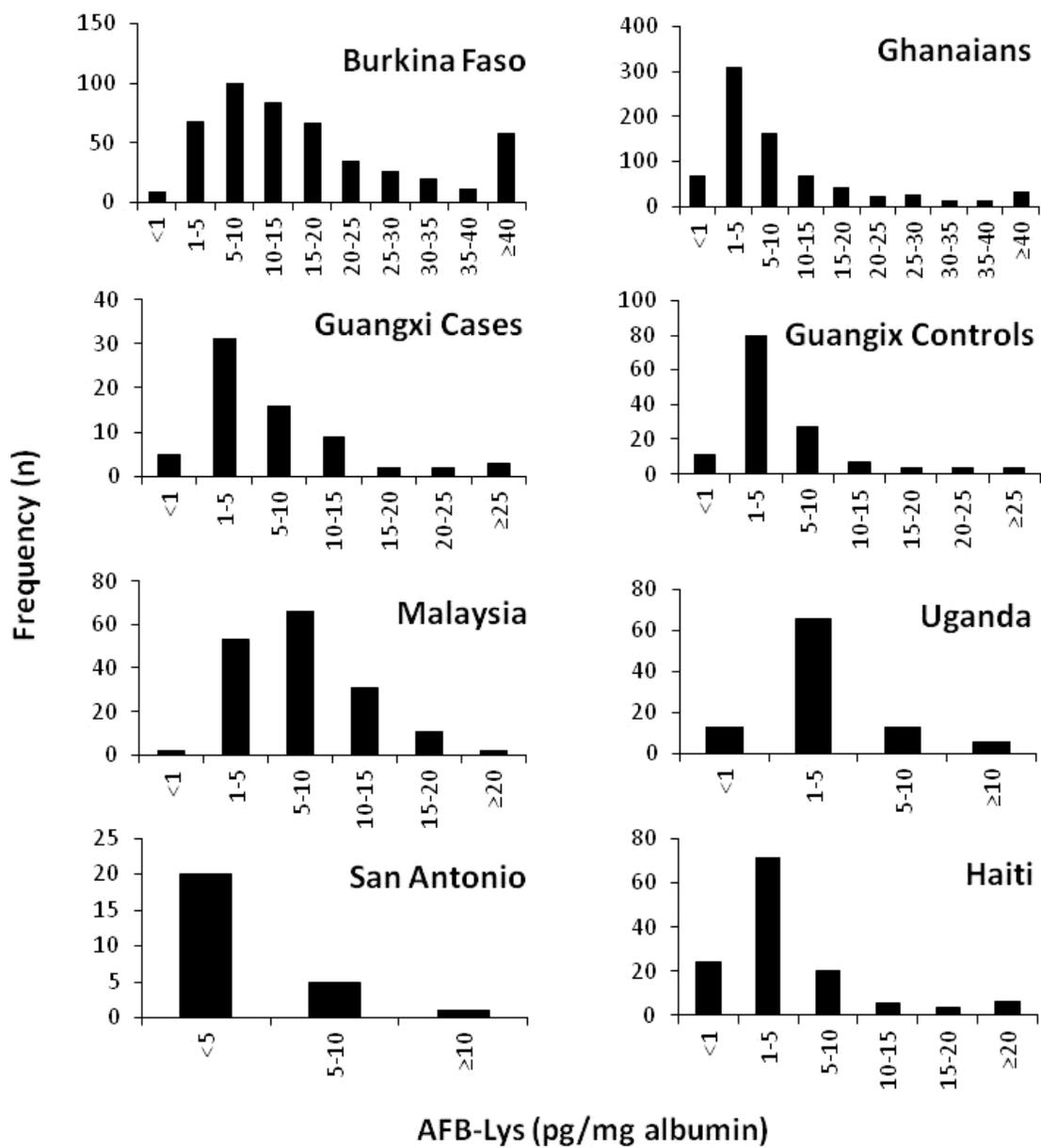


Figure 7.3. Distribution frequency of serum AFB-Lys levels in different study populations.

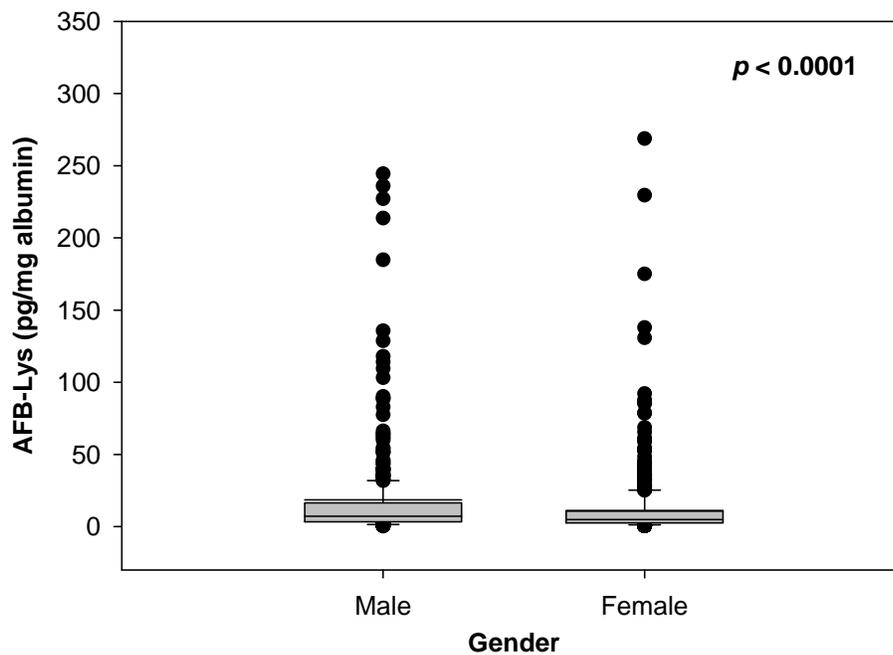


Figure 7.4. Overall gender distribution of serum AFB-Lys levels.

CHAPTER 8

CONCLUSION

This dissertation work was dedicated to the development of methods for detecting the human serum AFB-Lys adduct, a systematic evaluation of the toxicokinetics of this biomarker in rats, and the toxicity of AFB₁ in a sensitive rat model with special interest in the temporal histological, immunohistological, and the adverse immune effects. For the toxic effects evaluation, animal experiments that used both single-dose and repeated-dose approaches, were conducted. The development of a non-antibody AFB-Lys method facilitates assessment and comparison of human exposure to AFB₁ in many parts of the world, and serves as the first step towards the long-term goal of establishing a global network for studying global AFB₁ exposure and evaluating the efficacy of promising chemoprevention strategies. Overall, the advances and conclusions that have been accomplished by this dissertation work include:

1. Development and validation of a very sensitive solid-phase extraction based HPLC method with fluorescence detection. Among different test cartridges commercially available, the Water SPE MAXTM cartridge and Sep-PakTM cartridge showed better recovery than the HLBTM cartridge. The MAXTM cartridge is chosen for the method development because it is more time efficient than the Sep-PakTM cartridge. This non-antibody based method shows excellent sensitivity, with a detection limit of 0.4 pg/mg albumin. The inaccuracy and imprecision rates are less than 10% for intra- and inter-days analysis. This method is ready to be used in large-scale human serum samples analysis.

2. The single-dose treatment study in rats demonstrated that acute AFB₁ exposure produced a peak level of serum AFB-Lys at 4 h, followed by a rapid decrease. The pharmacokinetic analysis showed a half life of 2.31 days. Physiologically based pharmacologic model analysis showed that approximately 1.12-1.98% of AFB₁ dose is converted to serum AFB₁-Lys 24 h after exposure. Single-dose treatment elicited most pronounced histological changes (bile duct proliferation and periportal necrosis) and altered liver enzyme activities (e.g., serum ALT and AST) at 3 days after exposure.

3. Repeated-dose treatment in rats demonstrated that liver GST-P⁺ cells and foci are dose and time dependently induced, supporting the potential use of this marker as a surrogate endpoint in chemoprevention studies. Liver enzyme activity changes (ALT and AST) in serum were not as sensitive as the development of GST-P⁺ hepatocytes during repeated treatment. Repeated-dose study demonstrated that serum AFB-Lys levels linearly increases with time at low doses (5-25 µg/kg b.w.), indicative of the value of this biomarker to reflect chronic low dose exposure. Repeated AFB₁ exposure impaired the rat spleen lymphocyte function: short-term exposure (1-week) showed an overall inhibition of cytokine expression in CD4⁺, CD8⁺, and NK cells; long-term (5-week) exposure promoted IFN-γ and TNF-α secretion while suppressing IL-4 expression, suggesting the involvement of pro-inflammatory effects. The pro-inflammatory effects may not be reflective of histological changes and may imply other modes of action.

4. Human exposures to AFB₁ are highly variable among different populations and inter-individually. Populations from Sub-Saharan Africa and Southeast Asia areas were demonstrated to have much higher levels of serum AFB-Lys adduct and higher detection

rates compared to populations from the United States. The rank from high exposure to low is: sub-Sahara > Southeast Asia > Haiti > United States.

5. Collectively, our results demonstrate the validity of the non-antibody method in detecting the serum AFB₁-lysine adduct in both animal experiments and human populations.

CHAPTER 9

FUTURE DIRECTION

Aflatoxin research, started in the early 1960s, remains an active field of science at present. Through the literature, the trajectory of research on aflatoxins depicts intertwining lines of toxicological, agricultural, epidemiological, and regulatory efforts. There are good reasons for the continuing research on aflatoxins. The underlying economic cost, health effects and disease burdens, clinical implications, and mechanistic understanding of aflatoxin exposure cannot be ignored and stimulates research in many different fields.

Health costs and prevention

It has been estimated that worldwide, approximately 25% of crops are affected by mycotoxins annually ¹, of which aflatoxins are a major problem. How to prevent the great economic loss due to aflatoxin contamination of world crops remains a huge task for industry and regulatory agencies. The relative cost-effectiveness of aflatoxin reduction strategies in Africa have been assessed by Wu's group ^{2,3}. According to Wu's group ^{2,3}, both pre-harvest and post-harvest strategies have been proven cost-effective based on the World Health Organization (WHO) standard, and "public health interventions must be readily accepted by the public, and must have financial and infrastructural support to be feasible in the parts of the world where they are most needed". For the success of these public health intervention strategies (such as biocontrol, post-harvest package, NovaSil clay supplementation, and hepatitis B vaccination), public education, as well as financial and infrastructural support are critical especially in the less developed countries ⁴. Future

work will be directed to advance the application of these intervention approaches as well as to assess the efficacy employing molecular biomarkers in a diversity of populations which are at high risk for aflatoxin exposure.

Clinical implications of aflatoxin exposure

Besides the etiological role of aflatoxins in human hepatocellular carcinoma ⁵, aflatoxin exposure has been associated with several adverse health effects, including kwashiorkor ⁶, Reye syndrome ⁷, growth impairment ⁸, and immune suppression in children ⁹. Increasing evidence from epidemiological studies has suggested an association between aflatoxin exposure and those disease or adverse effects, however, a causal effect, if existing, warrants more specific studies especially in a prospective manner to verify. Still, other possible effects of continuous or intermittent dietary exposure to aflatoxins, which occurs widely in some developing countries, have received little study, such as the potential contribution of aflatoxin exposure to infectious diseases. Besides the well recognized interaction between aflatoxin and hepatitis B virus ^{5, 10-12}, other viruses such as tuberculosis and HIV may interplay with aflatoxins, considering the co-existence of both environmental agents in the tropical areas and circumstantial evidence from human studies that suggest such an effect ¹³⁻¹⁶. How to correlate the aflatoxin biomarkers with the risk or disease status remains a question to answer. To partially answer this question, the integration of data from clinical investigation and molecular biomarker measurement through the natural history of certain diseases might be useful to demonstrate such a link, which obviously will involve efforts from both doctors at the bed and scientists at the bench. Because most human diseases are an expression of the interaction between

individual genetic makeup and environmental agents ¹⁷, the individual phenotypes of the metabolic enzymes in particular, need to be known for an accurate risk analysis.

Mechanistic studies

Many human studies show an association between exposure to toxic agents and an increased risk of certain diseases. A mechanistic understanding of the relationship between exposure and disease depends on the availability of proper animal models and *in vitro* approaches. One difficulty associated with animal model development is the species difference in aflatoxin metabolism, which can sometimes be overcome by using transgenic animal models that specifically express human enzymes. Failure to demonstrate such an association in animal models sometimes may be related to the selection of endpoints, such as in the case of kwashiorkor ¹⁸. In that study, serum total protein and albumin, and histopathological changes were used to indicate the development of kwashiorkor. Lack of specific endpoints to the disease may prevent the accurate assessment of such a causal relationship. Development of disease specific biomarkers may help to dissect the relationship between disease and exposure to aflatoxins. It is worth noting that, the assumption that "aflatoxin causes diseases" is so well accepted that other possibilities may be conveniently ignored. For instance, the disease of kwashiorkor is largely a problem of "being ignored", as described by Williams ¹⁹, that is, a problem of malnutrition.

Being lipophilic compounds, aflatoxins can penetrate cell membrane by passive diffusion ²⁰. They can also penetrate by active uptake ²¹. AFB's lipophilicity favors its bioaccumulation through food chain. However, there is little, if any, evidence showing the concentration of aflatoxins in adipose tissues (the fast and efficient metabolism may

account for this). Several nutrition related diseases (such as kwashiorkor and environmental enteropathy) have been linked with aflatoxin exposure²². A common mechanism may involve the interference with the gut absorption, which can be mediated by affecting the normal gut microflora or increasing the permeability of gut membrane. The exact mechanisms deserve further research.

Interaction with or Confounding by Other Mycotoxins

Fumonisin are another group of mycotoxins that are frequently found in corn and animal feeds. Fumonisin B₁ (FB₁) has been shown to be associated with farm animal diseases and a potential etiological factor in human diseases²³. The co-occurrence of aflatoxins and fumonisins in corn has been a common finding worldwide^{24,25}. Interaction of AFB₁ and FB₁ in acute toxicity and carcinogenicity has been demonstrated in animal models²⁶⁻²⁸. William et al. has also found an association between HIV infection, the incidence of AIDS developing from those HIV infections and the inclusion of maize in the diets of Africans²⁹. This association, as noted by Brown³⁰, seems to be more closely related to the presence of fumonisins than aflatoxins. The potential interaction between AFB₁ and FB₁, and trichothecenes as well (e.g., T-2), in the etiology of human diseases remains largely unknown. Future work that directs risk assessment of mycotoxins should consider the co-existence of these mycotoxins.

In summary, efforts are needed to encourage the application of intervention strategies to reduce aflatoxin exposure and to advance a mechanistic understanding of the role of aflatoxin exposure in human health impairment. Integrative research incorporating both specific molecular biomarkers related to aflatoxin exposure and the adverse

outcomes is necessary if we are to develop a better mechanistic understanding of the role of aflatoxin in human diseases.

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