AN ABSTRACT OF THE THESIS OF

<u>Claudia McLaughlin Troxel</u> for the degree of <u>Doctor of Philosophy</u> in <u>Toxicology</u> presented on October 3, 1996. Title: <u>Aflatoxin B₁ Metabolism</u>, <u>CYP1A Induction</u>, and the Influence of CYP1A Induction on Aflatoxin B₁ Metabolism in Zebrafish.

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While small fish models are assuming prominence as alternative models for chemical carcinogenesis studies, little is known about the actual metabolism of carcinogens in these models. This thesis investigated the metabolism of the potent hepatocarcinogen aflatoxin B₁ (AFB₁) and its modulation by CYP1A induction in the zebrafish.

HPLC analysis showed that the major metabolites excreted into water following intraperitoneal injection of [³H]AFB₁ in zebrafish were aflatoxicol, aflatoxicol-glucuronide and unreacted AFB₁. Fish intraperitoneally injected with 50-400 μg [³H]AFB₁/kg body weight displayed a linear dose response for hepatic DNA-adduction at 24 hours. AFB₁-DNA adduct levels showed no statistical difference from 1 to 21 days after injection, suggesting poor adduct repair in this species. DNA binding in female fish was 1.7-fold higher than in males. An *in vitro* AFB₁ metabolism assay verified that zebrafish liver supports AFB₁ oxidation to the 8,9-epoxide proximate electrophile.

Investigations were conducted assessing CYP1A response in zebrafish. Intraperitoneal injection of 75-150 mg Aroclor 1254/kg body weight or dietary exposure to 500 ppm failed to induce CYP1A protein or associated 7-ethoxyresorufin-O-deethylase (EROD) activity. However, dietary β -naphthoflavone (β NF) at 500 ppm induced CYP1A approximately 3-fold above controls. A single intraperitoneal injection of 150 mg β NF/kg body weight showed maximum CYP1A protein levels between 24 to 36 hours, while EROD activity peaked before 48 hours. CYP1A showed dose-responsiveness in fish intraperitoneally injected with 25-150 mg β NF/kg body weight. Dietary exposure to 0.75 ppm 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) significantly induced CYP1A.

The effect of TCDD on the metabolism of AFB₁ in zebrafish was examined. In addition to aflatoxicol, aflatoxicol-glucuronide, and parent AFB₁ excreted by the control AFB₁ group, the TCDD-pretreated group also produced unresolved aflatoxin M₁/aflatoxicol-M₁ and aflatoxicol-M₁-glucuronide. Hepatic AFB₁-DNA adduction was approximately 4-fold higher in the TCDD treated group compared to controls. This significant difference could not be explained by increased capacity for bioactivation of AFB₁ as measured by an *in vitro* AFB₁ metabolism assay. However, it was demonstrated that zebrafish have the capacity to bioactivate aflatoxin M₁ to a reactive intermediate, and it is proposed that this intermediate may be responsible for the increased DNA binding.

AFLATOXIN B₁ METABOLISM, CYP1A INDUCTION, AND THE INFLUENCE OF CYP1A INDUCTION ON AFLATOXIN B₁ METABOLISM IN ZEBRAFISH

by

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AFLATOXIN B₁ METABOLISM, CYP1A INDUCTION, AND THE INFLUENCE OF CYP1A INDUCTION ON AFLATOXIN B₁ METABOLISM IN ZEBRAFISH

CHAPTER 1

INTRODUCTION

Fish are becoming an increasingly important model in science. While the first studies investigating metabolism in fish suggested low or no biotransformation capabilities, subsequent investigations have shown that is not the case. Not only do fish possess many of the metabolizing enzymes that mammals do, but they also are susceptible to the carcinogenic process. Fish were found to be responsive to carcinogens when an epizootic outbreak of liver tumors developed in hatchery-reared trout. The carcinogen responsible for the tumors was identified as the potent mycotoxin aflatoxin B₁ (AFB₁), which was a contaminant of the diet. The rainbow trout has since been shown to be the most sensitive species to aflatoxin's carcinogenic effects (Bailey et al., 1996). Tumors have also been found in feral fish populations, particularly bottom feeding fish. For many of these cancer epizootics, correlations have been made between the presence of neoplasia and environmental contamination, especially in bottom-feeding fish (Moore and Myers, 1994; Hawkins et al., 1995). For example, polycyclic aromatic hydrocarbons (PAHs) have been linked to hepatic lesions in English sole from Puget Sound and brown bullhead in the Black River (Hawkins et al., 1995, and references therein). It was also discovered that when fish are exposed to some of these compounds, such as the PAHs and polychlorinated biphenyls (PCBs), a particular enzyme, cytochrome P4501A, becomes induced, and can serve as a marker of recent exposure.

Many species of small fish are currently being developed as models for carcinogenesis. Rainbow trout have certainly proven to be a valuable model for carcinogenesis studies, but trout are limited by several factors including an annual spawning cycle, delayed maturity until approximately 2 years of age, and the requirement of cold water temperatures such that they are unsuitable for many *in situ* exposure protocols. Most aquarium fish, on the other hand, spawn regularly, reach maturity within months, and can withstand a wide range of temperatures. The guppy, Japanese medaka, and fathead minnow are only a few of the small fish species being investigated. While numerous studies have been and continue to be conducted investigating the susceptibility of these various species of fish to carcinogens, surprisingly few studies have explored the mechanistic basis behind the results of these exposures.

My thesis work was part of a U.S. ARMY grant assessing the suitability of zebrafish as a model for carcinogenesis. The Japanese medaka is currently the small fish of choice for these studies, but no one species is sensitive to all carcinogens. Therefore, zebrafish are being investigated for their suitability as a complimentary species. Zebrafish have the added advantage of being a popular model for the study of genetics and developmental biology. Chapter two explores *in vivo* aflatoxin B₁ metabolism and hepatic DNA-adduction after intraperitoneal administration. The experiments described therein investigate the capacity of the zebrafish for bioactivation

and detoxification of this carcinogen. It has been shown by other studies conducted in our laboratory that zebrafish are resistant to the carcinogenic effects of AFB₁ when this mycotoxin is administered in the diet.

Environmental exposures are seldom limited to just one chemical or carcinogen as they are under the carefully controlled laboratory settings. Coexposures can have a variety of effects, ranging from protection to enhancement of the toxicity or carcinogenicity of a compound. As mentioned earlier, CYP1A is an enzyme commonly elevated when an animal has been exposed to any number of 3-methylcholanthrene-type inducers. The consequence of such induction in the normal biology of the animal is not certain. Chapter 3 characterizes the induction of CYP1A in zebrafish following exposure to three different inducers; the commercial PCB mixture Aroclor 1254, the synthetic flavone β -naphthoflavone (β NF), and 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD), the most potent inducer known. The consequence of induction of CYP1A and the other associated enzymes following dietary administration of TCDD on AFB₁ metabolism and hepatic DNA adduction is then addressed.

Chemical Carcinogenesis

One of the first documentations that an environmental agent is capable of causing cancer was presented by Sir Percival Pott in the late 1700's. He linked the high incidence of scrotal cancer in chimney sweeps to exposure to soot and coal tar. It was not until the early 1900s that Pott's hypothesis of the carcinogenic potential of the tar was tested and confirmed in a laboratory setting by Yamagiwa and Ichikawa, who found dermal application of coal tar on rabbits caused cancer (reviewed in Phillips, 1983).

Today, the American Cancer Society estimates that one in four individuals in America will develop cancer over the course of their lifetime. Doll and Peto (1981) proposed that approximately 80% of all human cancer cases in the United States are caused by dietary factors, tobacco smoke, and other environmental exposures. Because of the prominence and severity of this disease, much attention has been focused on cancer research.

While evidence of carcinogenicity in humans obtained by epidemiological studies is the most desirable, these data are not often available. Therefore, scientists have turned to laboratory animals for testing the potential of an agent to cause cancer. Rodent models have been preferred for these studies, but they have several limitations. For example, these experiments are time consuming and expensive, and therefore only a limited number of compounds can be tested per year. In addition, the mandatory use of the maximum tolerated dose in these studies calls into question the validity of the results (Ames and Gold, 1991; Ames *et al.*, 1993). Furthermore, many granting agencies are encouraging the supplementation of mammalian models with alternative models and approaches.

For the past three decades, the rainbow trout model has been investigated and developed as a nonmammalian alternative model for carcinogenesis studies (reviewed by Bailey et al., 1996). The first demonstration of the susceptibility of rainbow trout to carcinogens occurred in the late 1950's and early 1960's, during an epizootic outbreak of liver tumors in hatchery-reared trout. The cause was traced to a contaminant in cottonseed oil used in the diet, which was identified as the potent mycotoxin, AFB₁ (Sinnhuber et al., 1968; Ayres, et al., 1971). Since then, rainbow

trout have proven to be susceptible to a number of other various carcinogens as well, including dimethylnitrosamine (Ashley and Halver, 1968; Grieco et al., 1978), diethylnitrosamine (DEN) (Shelton et al., 1984; Hendricks, et al., 1994) and benzo[a]pyrene (Hendricks et al., 1985).

Rainbow trout are not the only fish species susceptible to the carcinogenic process. In a hallmark study in 1965, Stanton demonstrated that DEN was capable of inducing hepatic tumors in the zebrafish. Further studies confirmed the responsiveness of zebrafish to DEN and also to nitrosomorpholine, although a water exposure to aflatoxin B₁ failed to elicit tumors (Pliss and Khudoley, 1975; Khudoley, 1984; Bauer et al. 1972). Surprisingly, the research on carcinogenesis in zebrafish ended with these few studies. Information on the sensitivity of many other small fish models is on the increase, however (reviewed in Hawkins et al., 1995).

Carcinogenic Process

Cancer is generally believed to be a multistage process involving initiation, promotion, and progression, although much debate surrounds the details associated with these stages. The following is only a brief summary of some of the theories concerning the possible mechanisms of carcinogenesis. Most of the basic principles can be found in Williams and Weisburger (1991), Weinberg (1989a), Pitot and Dragen (1991), and Harris (1991).

Initiation

The first step of the carcinogenic process is initiation, and usually involves electrophilic interaction of a compound with DNA. There are two classifications of initiators: direct-acting carcinogens and procarcinogens. Direct-acting carcinogens are reactive electrophiles requiring no further activation, such as many cytostatic agents used in chemotherapy. Mechanisms of action include alkylation and acylation of DNA. Procarcinogens, on the other hand, require metabolic activation to a reactive intermediate, or proximate carcinogen (reviewed by Guengerich, 1992). It is this electrophilic intermediate which can then bind to macromolecules including DNA (Miller and Miller, 1981). The DNA adduct resulting from the electrophile binding to DNA may be repaired, may lead to cell death if the damage is extensive, or may become a permanent DNA mutation following replication.

The consequence of mutagenesis depends on numerous factors, including the type and position of the mutation. Point mutations occur when a single base is substituted. When a purine is substituted for another purine, or a pyrimidine for a pyrimidine, the substitution is a transition; conversely, a transversion occurs when a purine is replaced by a pyrimidine (or visa versa). Consequences of point mutations range from having virtually no effect on gene function to completely inactivating a gene product. Frameshift mutations arise when a base is inserted or deleted, resulting in a shift in the reading frame of a gene. Usually the gene product is nonfunctional. Not only is the type of mutation important, but the location of the mutation on the DNA is also crucial. If it occurs in a non-critical portion of the gene, such as on a non-transcribed region, there may be no consequence. Mutations leading to the activation of oncogenes or the

inactivation of tumor suppressor genes, on the other hand, can have a substantial impact, which will be discussed in the following section. Once a gene containing a mutagenic DNA lesion undergoes a round of replication without repair, the mutation becomes fixed and the change is irreversible.

Promotion

The initiated cell can remain dormant until appropriate factors, such as hormones, viruses, or chemicals, stimulate cellular proliferation. This reversible process is called promotion, and permits the clonal expansion of cells to a preneoplastic lesion. Increased cellular proliferation also decreases the available amount of time for repairing DNA, possibly leading to genomic instability (Swenberg, 1993). Activation of oncogenes and/or inactivation of tumor suppressor genes (or antioncogenes) is believed to play a role in the transformation to neoplasia.

Proto-oncogenes are normal constituents of the cell that are important for cellular function, but when activated, appear to play a central role in the carcinogenesis process. Proto-oncogenes can be activated to oncogenes by mutations, gene amplification, or chromosomal translocations (reviewed in Ladik and Förner, 1994). Such activation can lead to a loss of growth regulation and differentiation of the cell (reviewed in Smith *et al.*, 1993). There is speculation that activation of at least two oncogenes is required for the full neoplastic conversion of a cell, although not all the evidence supports this model (Weinberg, 1989b). A family of oncogenes known as *ras* oncogenes are commonly mutated at codons 12, 13, and 61 in human tumors, but the percentage of *ras* mutations varies greatly depending on the tumor type (Bos, 1989). Importantly, *ras* mutations are

not limited to just human tumors, but have been found in tumors in many other species including fish (Chang et al., 1991; Van Beneden and Ostrander, 1994).

Tumor suppressor genes or antioncogenes are also essential for normal cellular function, and their numerous functions include regulating apoptosis and growth, DNA synthesis and repair, involvement in intercellular communication, and protease inhibition (Harris, 1991; Greenblatt *et al.*, 1994). In contrast to the oncogenes, inactivation of the tumor suppressor gene is required for cellular transformation, and requires the loss of function of both alleles. The *p53* suppressor gene is currently one of the most commonly mutated genes detected in human tumors, and unlike the *ras* mutations, *p53* mutations have been discovered in over 30 codons (Purchase, 1994; for review see Greenblatt *et al.*, 1994). However, there has often been a rather poor correlation between species in type and frequency of *ras* mutations and, in particular, *p53* mutations (McMahon, 1994; Eaton and Gallagher, 1994).

Progression

The irreversible transformation of benign, preneoplastic lesions to malignant neoplasms is termed progression. During this final stage of carcinogenesis, the neoplastic cells frequently invade surrounding tissues and organs and enter the lymphatic or vascular system to be carried to other sites in the body for implantation (metastasis). Malignant neoplasms are associated with karyotypic changes, lack of cellular differentiation, metastasis, increased growth rate, and a heterogenous cell population (Weinberg, 1989a; Robbins and Kumar, 1987). Activation and/or inactivation of critical genes is believed to play an important role in this final stage of carcinogenesis

as well, but there is still much to learn about which genes are actually involved. The future holds much promise in further elucidating more of the many important factors associated with this disease.

Biotransformation in Fish

Although it had been originally hypothesized that aquatic organisms were incapable of metabolizing xenobiotics, numerous studies have since proven the contrary. Indeed, biotransformation of xenobiotics is just as significant in aquatic species as it is in mammals.

Phase I Metabolism

Biotransformation is usually divided into two phases: phase I and phase II metabolism. Phase I metabolism serves to add or expose functional groups, thereby increasing the compound's polarity. While the increased polarity may allow the compound to be directly excreted, phase I metabolism typically primes the xenobiotic for conjugation by a phase II enzyme. The reactions involved in phase I metabolism include oxidation, reduction, and hydrolysis, and utilize a multitude of different enzymes (Sipes and Gandolfi, 1991; Hodgson and Levi, 1994).

Organic lipophilic compounds are metabolized primarily by oxidation via the mixed function oxidases (MFOs) and flavin monooxygenase (FMO). Although these enzyme systems are more thoroughly characterized in mammalian systems, research in aquatic species is growing. FMO metabolizes primary, secondary, and tertiary amines,

and sulfur-containing compounds (Sipes and Gandolfi, 1991; Ziegler, 1993). This membrane bound enzyme requires the electron donor NADPH and the prosthetic group FAD. In fish, FMOs are important in N-oxidation reactions, such as the oxidation of the endogenous substrate trimethylamine to trimethylamine oxide (Goldstein and Dewitt-Harley, 1973; Agustsson and Strom, 1981; Stegeman and Hahn, 1994). FMO-like activity has been detected in rainbow trout (Schlenk and Buhler, 1991a; 1991b; 1993), cod (Agustsson and Strom, 1981), striped bass (Cashman *et al.*, 1990), and nurse shark (Goldstein and Dewitt, 1973), although no FMO activity was detected in channel catfish (Schlenk *et al.*, 1993).

The cytochrome P450s, a superfamily of heme proteins, are the most prominent class of phase I enzymes. These enzymes are important in the metabolism of both endogenous and exogenous substrates, including fatty acids, steroids, prostaglandins, drugs, and carcinogens. The MFO system requires the utilization of two enzymes: the heme protein, cytochrome P450, and NADPH-cytochrome P450 reductase, a flavoprotein responsible for shuttling electrons to the P450 enzyme and requiring NADPH as a cofactor. Although only one NADPH cytochrome P450 reductase enzyme has been isolated, Nebert and Nelson (1991) predict that "each mammalian species might easily have at least 60 and perhaps more than 200 individual P450 genes". Currently, P450s have been found in every phylum investigated.

In fish livers, total amounts of P450s vary from 0.1 nmol/mg to 2.0 nmol/mg protein, although average hepatic levels in untreated fish range from 0.2 to 0.5 nmol/mg (Stegeman, 1989). Several P450s have been purified or cloned from various fish species. P450s identified in rainbow trout include members of families 1A, 2K, and a

number of P450s involved in steroid metabolism such as 11, 17, and 19 (reviewed in Stegeman and Hahn, 1994). One of the catalytic functions of CYP2K1 is the bioactivation of aflatoxin B₁ to the reactive intermediate, which is capable of binding to protein, DNA, and RNA (Williams and Buhler, 1983). Topminnows appear to have a P450 with properties similar to CYP2E1, including the ability to bioactivate the procarcinogen nitrosodiethylamine (Kaplan *et al.*, 1991). Scup appear to possess CYP1A, CYP2B and a CYP3A-like isozyme (Stegeman and Hahn, 1994; Husoy *et al.*, 1994), and studies in zebrafish show expression of CYP1A- and CYP2K1-like proteins (Buchmann *et al.*, 1993; Collodi *et al.*, 1994). CYP1A has also been detected in induced cod, perch, plaice, little skate, and medaka (Stegeman and Hahn, 1994; Schell *et al.*, 1987). Immunoblots from eight fish species show that the apparent molecular weight of hepatic CYP1A in fish ranges from 54,000-59,000 daltons (Goksoyr *et al.*, 1991; Collodi *et al.*, 1994).

CYP1A is readily induced by a number of compounds including PAHs (benzo[a]pyrene, benzanthracenes), many flavones and indoles, 3-methylcholanthrene, and halogenated aromatic hydrocarbons (biphenyls, dibenzo-p-dioxins and dibenzofurans). CYP1A response in fish is currently being developed as a biomarker for certain environmental exposures (Goksoyr, 1995; Stegeman and Lech, 1991; Wirgin et al., 1994). Levels of CYP1A can be assessed by measuring catalytic activity, antibody detection of protein, and cDNA probes for CYP1A mRNA. Because this P450 plays a critical role in the metabolism of aromatic hydrocarbon procarcinogens, much research is now devoted to unraveling the mechanism of induction of CYP1A and

understanding the many factors, such as temperature and reproductive status, which influence this response.

In mammals, the induction of CYP1A1 is regulated at the gene transcription level via the Ah (aryl hydrocarbon) receptor (AhR), a helix-loop-helix DNA-binding protein. In the unliganded state, the Ah receptor is found in the cytosol of the cell associated with two 90 kDa proteins (at least one is heat shock protein 90 (Hsp90)) and possibly a 46 kDa protein. Upon diffusion into the cell, the ligand binds to the Ah receptor. After ligand binding, the proteins dissociate from the complex, the receptor complex translocates to the nucleus, and the ARNT protein (Ah receptor nuclear translocator protein) binds to the transformed AhR. The order of these events and the cellular location of ARNT are currently unclear, and proteins other than Hsp90 and ARNT are believed to play a role in the process, such as proteins involved in phosphorylation/ dephosphorylation of ARNT and the Ah receptor. Once in the nucleus, the receptor binds certain regions in DNA, including the dioxin- or xenobiotic-responsive regulatory elements, (DREs, XREs) upstream of CYP1A1 in DNA, increasing the rate of transcription of this gene product (Ah receptor regulation reviewed in Hankinson, 1995; Whitlock, 1993; Okey et al., 1994). It is important to note that in mammals the Ah receptor is also responsible for upregulating other enzymes (such as glucuronosyl transferase, quinone-oxido-reductase, glutathione-S-transferase, CYP1A2, and CYP2B1) and for inducing proteins involved in cell growth and differentiation (Hankinson, 1995). Evidence from studies in fish show consistency of CYP1A induction with a receptormediated response, and there has been a correlation between induction and the presence of the Ah receptor in fish (Stegeman, 1993; Stegeman and Hahn, 1994).

Phase II Metabolism

Phase II enzymes typically conjugate phase I metabolites with an endogenous molecule, such as a sugar moiety, a sulfate group, an amino acid, or glutathione. The addition of this conjugate increases the water solubility of the metabolite even further, commonly leading to excretion via bile or the kidneys. While there is evidence of some of the other phase II reactions, such as sulfation, occurring in fish, the most prominent piscine phase II reactions identified and studied are glucuronidation and glutathione conjugation.

Glucuronidation is just as important in fish as in mammals in the metabolism of endogenous compounds, such as sex steroids and bilirubin, and xenobiotics containing an appropriate functional group (James, 1987; reviewed in Clarke et al., 1991; George, 1994). In general, uridine diphosphate glucuronosyltransferases (UDPGT) catalyze the transfer of glucuronic acid from the high energy cofactor UDP-glucuronic acid to the substrate to form O-glucuronides (phenols, alcohols, and carboxylic acids), N-glucuronides (carbamates, sulfonamides, and aromatic amines), and S- or C-glucuronides (thiols, thiocarbamic acids, some drugs) (Dauterman, 1994; Gibson and Skett, 1986). In fish, O-glucuronides are the most common conjugates, followed by N-glucuronides. As in mammals, UDPGT activity is microsomal. While primarily located in the liver, UDPGT activity is also found in the kidney, gill and intestines (George, 1994). Benzo[a]pyrene, aflatoxin B₁, benzene, and tetrachlorobiphenyl are just a few of the xenobiotics that can be glucuronidated by fish (Clarke et al., 1991).

Glutathione-S-transferases (GSTs) mediate the conjugation of the endogenous tripeptide, glutathione, with an endogenous or exogenous electrophilic metabolite. These ubiquitous, primarily cytosolic, enzymes exist as homo- or heterodimers. In mammals, the GSTs are currently grouped into four classes, and all but one class exhibit activity for the substrate 1-chloro-2,4-dinitrobenzene (CDNB) (George, 1994; Dauterman, 1994). Following conjugation with glutathione, the metabolite may be directly excreted, or more typically, further metabolized to a mercapturic acid and then excreted. All fish examined so far demonstrate activity for CDNB, and GSTs have been purified in several fish species (reviewed in George, 1994). It appears from these studies that fish also have multiple isoforms consisting of two subunits. Tissue-specific expression of the various GSTs appears to depend on both the specific isoform and fish species, but the GSTs are predominately found in liver, kidney, gills, and intestine (George, 1994).

Aflatoxin B, Metabolism and DNA Adduction

Aspergillus paraciticus. These mycotoxins frequently contaminate improperly stored foodstuffs, such as corn and peanuts. Aflatoxin B₁ (AFB₁) is the most potent metabolite produced, and has recently been classified as a Group I carcinogen in humans (IARC, 1993). Species response to this carcinogen is varied. Mice, coho salmon, and channel catfish are fairly resistant whereas rainbow trout, ducks and rats are susceptible to AFB₁-induced carcinogenesis. Investigations have shown that such species differences

in carcinogenic response can generally be explained by differences in biotransformation of this carcinogen (Eaton et al., 1994; Eaton and Gallagher, 1994).

Aflatoxin B, Metabolism

AFB₁ requires metabolic bioactivation to the reactive intermediate, the AFB₁-8,9-epoxide (AFBO), to exert its biological effects (Essigman *et al.*, 1982; Swenson *et al.*, 1977). It is this unstable metabolite that is capable of binding to RNA, DNA, and other cellular macromolecules. The bioactivation of AFB₁ is typically a P450-mediated reaction, although alternative pathways, such as prostaglandin H synthetase- and lipooxygenase- mediated pathways, do exist. The primary P450s responsible for bioactivation of AFB₁ in humans are CYP1A2 and CYP3A4 (Gallagher *et al.*, 1994; Shimada and Guengerich, 1989; Eaton and Gallagher, 1994). CYP2K1 catalyzes AFB₁ activation in rainbow trout, and this P450 displays higher activity towards AFBO formation than rat P450s (Williams and Buhler, 1983).

Differences in the production of the epoxide can sometimes explain the variations in sensitivity between species, such as rainbow trout compared to rat and Coho salmon (Williams and Buhler, 1983; Bailey et al., 1988). However, phase II detoxification reactions also can play a crucial role. Although mice actually have a higher microsomal activity for AFBO production compared to rats, mice are resistant to AFB₁ carcinogenicity due to a high activity of glutathione-S-transferase toward the epoxide (Degen and Neuman, 1981; Eaton et al., 1994; Eaton and Ramsdell, 1992). Glutathione conjugation of AFBO does not appear to be a significant reaction in fish (Gallagher and Eaton, 1995; Valsta et al., 1988).

Other metabolic pathways include additional phase I and phase II reactions. Reduction of AFB, by a cytosolic reductase generates aflatoxicol (AFL), a metabolite identified in rats, rabbits, birds, and fish (Eaton et al., 1994; Gallagher and Eaton, 1995; Toledo et al., 1987). In mammals, O-demethylation or hydroxylation produces the metabolites aflatoxin P_1 (AFP₁), aflatoxin Q_1 (AFQ₁), and aflatoxin M_1 (AFM₁). Trout can also form AFM, and can further reduce it to aflatoxicol M, (AFL-M₁) (Loveland et al., 1979; 1983). AFL and AFL-M₁ can be further metabolized by glucuronidation, a particularly important reaction in fish (Loveland et al., 1984; Toledo et al., 1987). While hydroxylated metabolites of AFB, are generally considered to be detoxification products, the metabolites AFM₁ and AFL are still carcinogenic (Eaton et al., 1994; Shoenhard et al., 1981; Nixon et al., 1981; Bailey et al., 1994). In addition, in some species, AFL serves as a reservoir for AFB, in that it can be oxidized back to AFB, for subsequent bioactivation (Salhab and Edwards, 1977; Loveland et al., 1977). A summary of the AFB₁ biotransformation pathways in rainbow trout is found in Figure 1.1.

Significance of aflatoxin B_1 DNA adducts

Aflatoxin-8,9-epoxide, the reactive intermediate produced by bioactivation of AFB₁, can bind to cellular DNA, primarily forming the *trans*-8,9-dihydro-8-(N⁷-guanyl)-9-hydroxyaflatoxin B₁ adduct (Swenson *et al.*, 1977; Eaton and Gallagher, 1994; Bailey, 1994). This adduct can undergo rearrangement to a ring opened formamidopyrimidine (FAPY) derivative, which is very persistent over time. Despite its persistence, however, evidence points to the initial adduct formed being the most important in

tumorigenesis (Bailey, 1994). Molecular dosimetry studies in both rats and trout demonstrate that DNA adduct formation highly correlates with tumor response, as does AFB₁ dose with the number of DNA adducts generated. These correlations are even more powerful because they display a strong linear relationship, even at low doses, suggesting that there is not a threshold for AFB₁ genotoxicity (Bechtel, 1989; Appleton et al., 1982; Choy, 1993; Dashwood et al., 1992; Bailey, 1994).

Figure 1.1 Biotransformation pathways of AFB₁ in rainbow trout. Abbreviations used: AFM₁, aflatoxin M_1 ; AFL- M_1 , aflatoxin M_1 ; AFL, aflatoxicol; UDPGT, UDP glucuronosyltransferase; GST, glutathione-S-transferase (from Breinholt, 1994).

The location of AFB₁-induced mutagenesis again depends on the species and gene under investigation. *Ras* mutations, and in particular Ki-*ras* mutations, have been identified in both rat and trout exposed to AFB₁, with hotspots at codons 12 and 13 (Bailey, 1994; Eaton and Gallagher, 1994). Studies in humans have yet to detect activated oncogenes in populations exposed to AFB₁. It appears that inactivation of the *p53* tumor suppressor gene plays a role in humans. An association has been found between mutation in codon 249 in *p53* and liver tumors from individuals from aflatoxin-endemic areas (Harris, 1991). Further refinements in the sensitivity of techniques used in detecting mutations should help to elucidate other molecular events critical in AFB₁-induced carcinogenesis.

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CHAPTER 2

IN VIVO AFLATOXIN B₁ METABOLISM AND HEPATIC DNA ADDUCTION IN ZEBRAFISH (Danio rerio)

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Abstract

The zebrafish (Danio rerio) is assuming prominence in developmental genetics research. By comparison, little is known of tumorigenesis and nothing is known of carcinogen metabolism in this species. This study evaluated the ability of zebrafish to metabolize a well-characterized human carcinogen, aflatoxin B₁ (AFB₁), to phase I and phase II metabolites and assessed hepatic AFB₁-DNA adduction in vivo. Fish i.p. injected with 50-400 µg [3H]AFB₁/kg body weight displayed a linear dose-response for hepatic DNA binding at 24 hours. AFB₁-DNA adduct levels among treatments showed no statistical difference over the period from 1 to 21 days after injection, suggesting poor adduct repair in this species. DNA binding in female fish was 1.7-fold higher than in males (p < 0.01). An in vitro AFB₁ metabolism assay verified that zebrafish liver extracts oxidize AFB₁ to the 8,9-epoxide proximate electrophile (Km = 79.0 ± 16.4 μ M, Vmax = 11.7 ± 1.4 pmol/min/mg protein at 28°C). The excretion of AFB₁ and its metabolites was also examined by HPLC. As is typical of other fish studied, major metabolites excreted were aflatoxicol (AFL) and aflatoxicol-glucuronide (AFL-g), followed by unreacted AFB₁. AFL appeared as early as 5 minutes after injection, whereas AFL-g was a significant metabolite after 18 hours. This study shows that in vivo administration of AFB₁ in zebrafish results in moderate adduction of the carcinogen to liver DNA, and that zebrafish have the capacity for both phase I and phase II metabolism of AFB₁. The approximate 4-fold difference between rainbow trout and zebrafish AFB1-DNA covalent binding index appears insufficient to explain the relative resistance of zebrafish to dietary AFB1 hepatocarcinogenicity.

Introduction

Aquarium fish are gaining use as alternative models for chemical carcinogenesis studies. Many contributing factors include short life span, short time to tumor response, ability to spawn regularly, reduced husbandry costs, the ability to include the whole fish sagittal-section on one microscope slide for histology, and a low spontaneous tumor incidence (Hawkins *et al.*, 1988). The zebrafish was the first aquarium species in which chemically-induced tumors were demonstrated (Stanton, 1965). Later studies confirmed their responsiveness to diethylnitrosamine and also to nitrosomorpholine (Khudoley, 1984; Pliss and Khudoley, 1975). However, subsequent investigations have focused on life stage effects and toxicity upon exposure to various chemicals and heavy metals (Braunbeck *et al.*, 1990a; 1990b; Nagel *et al.*, 1991, Kalsch *et al.*, 1991; Bresch *et al.*, 1990; Nielson *et al.*, 1990; Dave and Xiu, 1991), with surprisingly little work on the response of zebrafish to carcinogens.

Aflatoxins are secondary metabolites produced by the fungus Aspergillis flavus, with aflatoxin B₁ identified as the most toxic metabolite. AFB₁ is a potent hepatocarcinogen and hepatotoxin in some species, such as the rat, duck, and rainbow trout, whereas other species, such as the mouse, seem to be fairly resistant to its effects. AFB₁ has recently been classified as a Group 1 carcinogen in humans (IARC, 1993). Aflatoxin exposure may occur via contaminated food, such as moldy corn and peanuts. Research has indicated that interspecies differences in sensitivity to this hepatocarcinogen can often be explained by variations in the biotransformation of AFB₁ (Eaton and Gallagher, 1994; Bechtel, 1989; Bailey et al., 1996). The activation of AFB₁ to the exo-AFB₁-8,9-epoxide is thought to be responsible for its carcinogenic

effects because this unstable, highly reactive intermediate can bind to cellular macromolecules including DNA (Essigman et al., 1982; Swenson et al., 1977). Recent studies have shown that the endo-AFB₁ epoxide stereoisomer can also be produced, and although this epoxide is the more stable stereoisomer, it does not appear to be as efficient in forming DNA adducts (Raney et al., 1992a; 1992b; Eaton and Gallagher, 1994). The rate and ratio of formation of the exo- and endo-AFB₁ epoxide stereoisomers depends on the species under investigation. Formation of this intermediate alone is not enough to account for carcinogenesis, since detoxification is possible via a glutathione-S-transferase mediated reaction. Successful scavenging of the exo-AFB₁ epoxide by this pathway has been shown to be responsible for lower susceptibility to the carcinogenic effects of AFB₁ in mouse compared to rat (Degen and Neumann, 1981; Raney et al., 1992a).

AFB₁ response in fish depends upon the species being investigated. For example, coho salmon and catfish are resistant but rainbow trout are one of the most sensitive species known to the carcinogenic effects of AFB₁ (reviewed in Hendricks, 1994; Bailey *et al.*, 1996). A study in 1968 showed that exposure to microgram quantities of AFB₁ was acutely toxic to zebrafish embryos and larvae (Abedi and McKinley, 1968). However, zebrafish did not develop tumors after 2 ppm water exposure for 3 days (Bauer *et. al.*, 1972). We have recently found AFB₁ to be a relatively weak hepatocarcinogen in this species (unpublished results). The purpose of this study was to characterize the *in vivo* metabolism and hepatic DNA adduction of AFB₁ in the zebrafish as a first step in assessing the mechanisms of AFB₁ resistance in this species.

Materials and Methods

Animals

Adult (sexually mature) male and female zebrafish with an average body weight of 0.5 g were obtained from the Food Toxicology and Nutrition Laboratory at Oregon State University. The zebrafish were maintained in 29 gallon aerated aquariums and a controlled temperature of 26°C (± 1) and a 14 hour light:10 hour dark photoperiod. During the week, fish were fed twice daily with TetraMin Staple Food (flakes) each morning, and a combination of Oregon Test Diet (OTD) (Sinnhuber et al., 1977) and brine shrimp in the afternoon on Monday, Wednesday, and Friday, and TetraMin Staple Food and brine shrimp on Tuesday and Thursday afternoons. On weekends, fish received only one feeding per day consisting of OTD and brine shrimp. Fingerling rainbow trout (Oncorhynchus mykiss) were reared in the Food Toxicology and Nutrition Laboratory as described (Sinnhuber et al., 1977).

Chemicals

AFB₁ was purchased from Sigma Chemical Company (St. Louis, MO), and was assessed for purity by thin layer chromatography (TLC); [³H]AFB₁ was from Moravek Biochemicals (Brea, CA) and the chemical and radiopurity was checked by TLC followed by radioscanning; Proteinase K, and RNase, DNase free were from Boehringer Mannheim Biochemicals (Indianapolis, IN); HPLC grade acetonitrile, methanol, tetrahydrofuran, and J.T. Baker C₁₈ Empore extraction disks were from VWR (Seattle,

WA); Hoechst #33258 from Calbiochem-Behring Corp. (La Jolla, CA); aflatoxin HPLC standards aflatoxicol (AFL), aflatoxicol-glucuronide (AFL-g), aflatoxin-M₁ (AFM₁), and aflatoxicol-M₁ (AFL-M₁) were prepared in our laboratory by previously published methods (Loveland et. al., 1983; 1984); all other chemicals were purchased from Sigma (St. Louis, MO) or Aldrich Chemical Company (Milwaukee, WI).

Metabolism of [H]AFB1

Adult female zebrafish, fasted for 24 hours, were intraperitoneally (i.p.) injected with 400 µg/kg body weight of [3H]AFB, (11.6 Ci/mmol) in 50% ethanol. The fish were rinsed with 2 ml of water after injection to account to remove residual traces of radioactivity, and placed in the dark in individual beakers containing 50 ml water. Water samples were collected and counted at various time points up to 24 hours for metabolite analysis. AFB, and metabolites were extracted from the aqueous samples using C₁₈ Empore Extraction Disks. The disks were washed with 45 ml of the elution solvent (10 mM potassium acetate (KOAc), pH 5.0, adjusted to 60% methanol (MeOH)) in 15 ml increments, and then conditioned with 45 ml of 100% MeOH, 45 ml of 50% MeOH, and 45 ml of 10% MeOH. Pooled water samples from each time point (approximately 150 ml) were adjusted to 10% methanol and were loaded onto the conditioned disks. Prior to elution, the disks were washed with 10 ml of 10 mM KOAc, pH 5.0, adjusted to 10% MeOH, to wash off proteins, salts, and exchanged tritium. AFB, and metabolites were eluted with 15 ml of 10 mM KOAc, pH 5.0, adjusted to 60% methanol. Total recovery rates (comprised of all fractions and the disk, which was cut up and counted) were greater than 85%, and the amount of radioactivity associated with the 60% methanol eluant represented 60-88% of the total radioactivity. The methanol eluates, containing the aflatoxin metabolites, were concentrated using a rotary evaporator and redissolved in 15% acetonitrile: methanol: tetrahydrofuran (AMT; 15:20:6 v:v:v) and 85% 0.02 M KOAc, pH 5.0. Reverse phase high pressure liquid chromatography was performed with a Beckman Model 334 instrument using a 4.6 x 150 mm, 5 micron Phenomenex C₁₈ column. The solvent system consisted of AMT (15:20:6 v:v:v) and 0.02 M KOAc, pH 5.0, with a 15 minute linear gradient from 15% AMT to 46% AMT. The flow rate was 1 ml/min, and UV detection was at 345 nm. Effluent samples of 30 drops (0.3 ml) were collected for liquid scintillation counting using a Beckman LS 7500 scintillation counter.

Hepatic aflatoxin-DNA adduction

To evaluate the dose-response of [³H]AFB₁-DNA binding, adult male and female zebrafish fasted for 24 hours were i.p. injected with 0, 50, 100, 200, or 400 μg/kg body weight of [³H]AFB₁ (11.6 Ci/mmol) in 50% EtOH and were rinsed with 2 ml of water to remove any residual skin contamination. Twenty-four hours after injection, livers were sampled with the gall bladders carefully removed, with 3 pools of 3 livers per dose. After sampling, the livers were not rinsed but immediately frozen in liquid nitrogen. DNA isolation and purification was based on the procedure by Strauss (1991), with the following modifications. Liver pieces were homogenized with digestion buffer. The samples were first extracted with equal volumes of phenol saturated with Tris-HCl buffer, pH 8.0., and then extracted twice with equal volumes of chloroform: isoamyl alcohol (24:1). The DNA was precipitated using 0.1 volume of 3 M sodium acetate and

2 volumes of cold 100% EtOH. After centrifugation, the EtOH was removed and the DNA allowed to dry. The DNA was dissolved in 0.5 ml of purified water by heating the samples at 38°C for 30 minutes, and then DNase-free RNase was added to the sample (final concentration $5\mu g/ml$), and the incubation continued for another 30 minutes. The samples were extracted twice with equal volumes of chloroform: isoamyl alcohol (24:1), and the DNA was again precipitated. After centrifugation, the EtOH was removed, and the DNA was washed with 70% ice-cold EtOH. DNA samples were stored at -20°C after dissolving the DNA in Tris-ethylenediaminetetraacetic acid (EDTA) (TE) buffer, pH 8.0. DNA concentration was quantified using the microfluorometric procedure described by Cesarone et al. (1979), with a Hoefer TKO100 DNA fluorometer. To determine the amount of [3H]AFB, bound to the DNA, samples were hydrolyzed by mixing with equal volumes of 1.0 M perchloric acid and heating at 70°C for 20 minutes. The hydrolyzed samples were then counted with a scintillation counter. Control liver samples spiked with [3H]AFB, during the DNA isolation process revealed that there was negligible carry-over of non-bound tritium through the isolation process.

To verify the sex-related difference in DNA binding observed in the dose-response study, an experiment was conducted with 9 groups of 3 females, and 7 groups of 3 males. The fish were i.p. injected with 400 μ g/kg body weight of [3 H]AFB₁ (13.0 Ci/mmol), the highest dose used in the dose-response experiment. Livers were sampled 24 hours later, and the amount of hepatic DNA binding was determined.

To evaluate the time-course of AFB₁-DNA binding, adult male and female zebrafish were i.p. injected with 400 μ g/kg body weight of [3 H]AFB₁ (15.4 Ci/mmol),

and livers were sampled 1, 2, 4, 7, 14, and 21 days after injection. Livers were immediately frozen in liquid nitrogen. DNA binding was determined as described above.

In vitro AFB₁-metabolism assay

The *in vitro* metabolism of AFB₁ was carried out with juvenile trout (n=10), or adult female (n=20) or male (n=6) zebrafish liver homogenates. The fish were fasted for 24 hours before sampling. Female zebrafish and juvenile trout livers were excised and immediately frozen in liquid nitrogen and stored at -80°C. Male zebrafish livers were immediately placed into homogenization buffer and homogenized and then frozen due to difficulties with protease release during liver excision. Livers were homogenized in a buffer containing 0.1 M potassium phosphate (pH 7.25), 20% glycerol, 1 mM EDTA, 0.1 mM butylated hydroxytoluene, 1 mM dithiolthreitol, and 0.1 mM phenylmethylsulfonylfluoride on ice. Samples were centrifuged at 1,000x g for 10 minutes, and the supernatant decanted and stored at -80°C. Protein concentration was determined by the method of Lowry *et al.* (1951), with bovine serum albumin as the standard.

The *in vitro* AFB₁-metabolism assay was based on the method of Monroe and Eaton (1987), as modified by Takahashi *et al.* (1996). Samples were preincubated for 2 minutes at room temperature before initiating the reaction with NADPH and reduced glutathione. Incubations were carried out for 45 minutes. Michaelis-Menton kinetics were determined with trout homogenates at both 13°C and 28°C using substrate (AFB₁) concentrations of 5, 10, 20 and 40 μ M AFB₁, and with female zebrafish homogenates

at 28°C at substrate concentrations of 10, 20, 40 and 80 μ M AFB₁. The initial rates of epoxide formation were linear with substrate concentration. Male zebrafish homogenates were assayed using 80 μ M AFB₁, the highest substrate concentration, for comparison to female zebrafish. All assays were conducted in duplicate. Incubations without homogenates were used to correct for the low inherent epoxidation activity of butylated hydroxyanisole (BHA)-induced mouse cytosolic protein included in the assay mixture.

Statistical analysis

Statistical analysis was performed with SAS, version 6.8 (SAS Institute Inc., 1989). Analysis was performed by either a one-way or two-way analysis of variance. A p value of less than 0.05 was considered significant.

Results

In vivo metabolism of [H]AFB₁

Early excretion kinetics of [³H]AFB₁ were evaluated by measuring the amount of radioactivity recovered in water within a 24 hour time period after i.p. injection (Figure 2.1). By 24 hours, 47% of the radioactivity administered to the zebrafish was recovered in the water. Excretion occurred rapidly with 25% of the administered radioactivity recovered by 45 minutes. This amount represented more than half of the total radioactivity recovered in the 24 hour time period. The remaining 53% of the

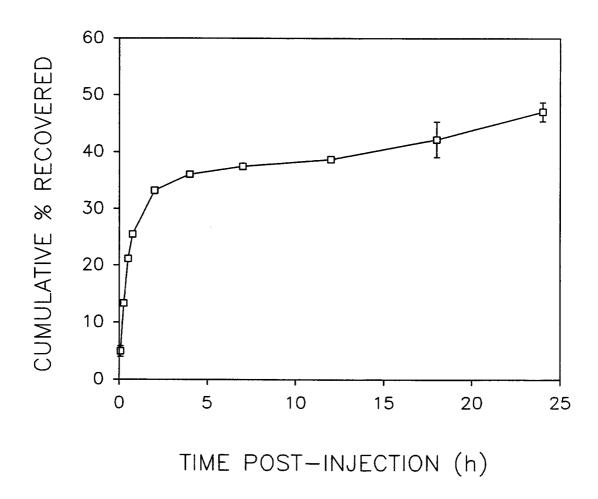


Figure 2.1. The cumulative percentage of radioactivity excreted into water within 24 hours after i.p. injection of $[^3H]AFB_1$. (n=3 females, error bars are \pm SEM).

administered dose is presumed to represent material found bound strongly to macromolecules and material eventually eliminated via fecal excretion.

The major metabolites of AFB₁ recovered in the water at various time points after i.p. injection were identified as aflatoxicol (AFL), aflatoxicol-glucuronide (AFL-g), and unreacted parent AFB₁ (Figure 2.2). Identification of these peaks was based on the retention times of known aflatoxin standards generated in our lab. Formation and excretion of AFL occurred rapidly in the zebrafish and was present in the water at 5 minutes, the earliest time point examined. Over the 24 hour time-period, AFL was the predominant metabolite, accounting for 17.3% of the original dose of AFB₁ administered. AFL-g and unreacted AFB₁ accounted for 4.1% and 3.6%, respectively.

Figure 2.3 shows the proportion of AFB₁ and metabolites recovered in the HPLC profile at each time point investigated, with values normalized to 100. AFL was the major metabolite recovered in the water at each time point up to 18 hours, accounting for 60 to 80% of the metabolites recovered. After 18 hours, AFL-g was the major metabolite, and represented over 80% of metabolites recovered at the 24 hour time point. An unidentified polar peak accounting for only a minor percentage of the metabolites was occasionally detected, being most prominent at 5 minutes. While this peak could represent a more polar metabolite, it did not co-elute with authentic AFB₁-glutathione conjugate, and was not reproducible in another experiment conducted investigating aflatoxin metabolism (Chapter 3, this thesis). The role of sulfate conjugation in AFB₁ metabolism by zebrafish investigated in another experiment by aryl sulfatase treatment indicated non-detectable levels of this conjugate (Chapter 3, this thesis). Unresolved aflatoxicol-M₁ (AFL-M₁) plus aflatoxin-M₁ (AFM₁) represented

only a very minor fraction of the metabolites, comprising less than 2% of the metabolites recovered at any one time point.

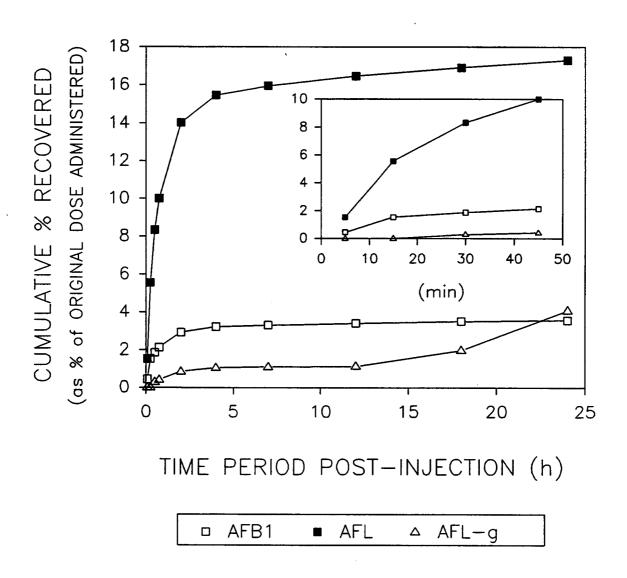


Figure 2.2. The cumulative percentage of AFB₁ and metabolites excreted into water as determined by HPLC as a percentage of the original dose administered. Each data point represents the pooled water samples from 3 females.

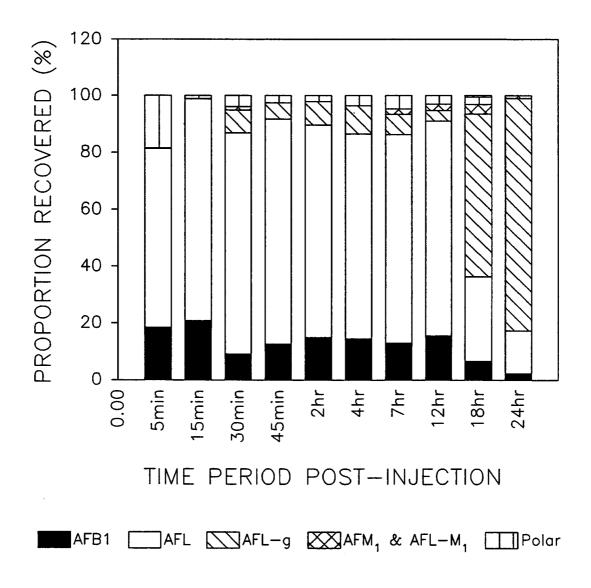


Figure 2.3. The proportion of AFB₁ and metabolites excreted into water at various time points over 24 hours as determined by HPLC. Values are normalized to 100.

In vivo AFB₁-DNA adduction in liver

The *in vivo* dose-response for AFB₁-DNA adduct formation was linear from 50-400 μ g AFB₁/kg body weight (Figure 2.4). Females had a higher level of DNA binding than the males at all doses tested. The covalent binding index (CBI), where CBI = μ mol chemical bound/mol DNA/mmol chemical administered/kg body weight, was 32,700 \pm 21,800 for males, and 56,000 \pm 15,500 for females. Females had a DNA binding index which was significantly higher (1.7 fold higher) than males (p < 0.01). A follow-up study investigating sex-related difference in AFB₁-DNA adduction verified the results found in the dose-response experiment. The binding index in the second experiment was 68,000 \pm 31,400 for females and 29,100 \pm 12,600 for males, which represents a 2.3 fold higher binding index in females compared to males (p < 0.05).

An investigation of the time-course of [³H]AFB₁-DNA binding was conducted to assess when maximum DNA adduct formation occurred, and to investigate the persistence of DNA adducts in this species (Figure 2.5). Although there was some trend toward highest adduction at day 7, the time-course experiment failed to show any statistical difference in binding levels from 1 to 21 days. Therefore, the time at which maximal adduct formation occurs could not be calculated. The half-life of the DNA adducts was also not evident from this experiment. However, a significant difference was again noted between male and female AFB₁-DNA binding levels, with the females exhibiting a binding index on average 42,000 units higher than the males (p < 0.001 when comparing the average of females to the average of males).

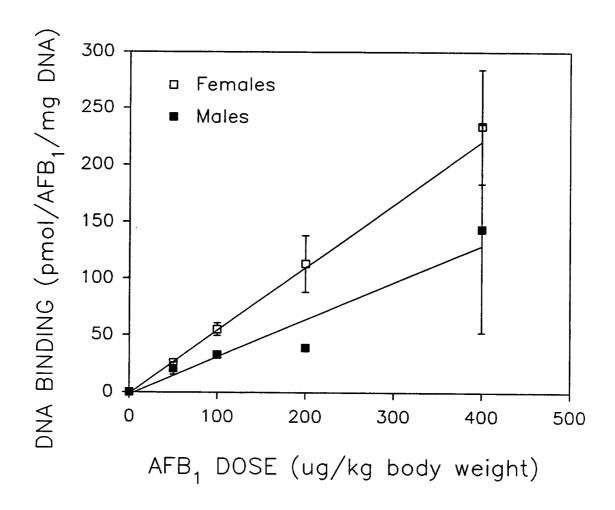


Figure 2.4. The *in vivo* dose response of hepatic aflatoxin-DNA adduct formation at 24 hours. Fish were i.p. injected with 50-400 μ g [3 H]AFB₁/kg body weight (data are means \pm SEM from 3 pools of 3 fish each.)

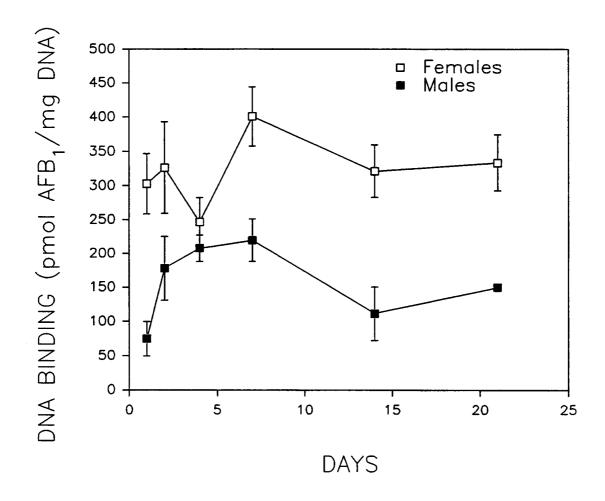


Figure 2.5. Aflatoxin-DNA adduct formation from 1-21 days. (Data are means \pm SEM from 3 pools of 3 fish each; data for males at Day 21 are mean \pm SEM from 2 pools of 3 fish each).

In vitro AFB₁-metabolism assay

Trout liver homogenates exhibited a higher Vmax for oxidation of AFB₁ to the 8,9-epoxide at both 13°C and 28°C compared to the female zebrafish. As was expected, trout Vmax decreased at the lower temperature. However, the values at 13°C are probably more representative of the physiological response in the trout, since this is their normal environmental temperature. At 13°C, the trout liver homogenates had a 3.1 fold higher Vmax and a similar Km when compared to female zebrafish homogenates at 28°C (Table 2.1) in this experiment. The 7-fold lower Km for AFL production suggests that the reaction would predominate over AFB₁-8-9-epoxide formation at low AFB₁ concentrations *in vivo* in zebrafish, and is consistent with the rapid excretion of AFL within 5 minutes of AFB₁ treatment.

Not surprising was the relatively high amount of aflatoxicol which was formed in the zebrafish liver homogenates. At 80 μ M AFB₁, male and female liver homogenates exhibited comparable activities for aflatoxicol production, with the female zebrafish liver homogenates possessing an activity of 8.17 \pm 0.09 pmol/min/mg protein, and male zebrafish an activity of 8.70 \pm 0.04 pmol/min/mg protein. Male zebrafish did appear to have a lower activity towards formation of the AFB₁-epoxide than females (4.10 \pm 0.08 pmol/min/mg protein compared to 5.90 \pm 0.10 pmol/min/mg protein, p < 0.01).

Table 2.1. Michaelis-Menton kinetics of AFB₁-epoxide formation in rainbow trout and zebrafish liver homogenates and aflatoxicol formation in zebrafish liver homogenates with the substrate AFB₁*.

Species	Temperature	Km (μM)	Vmax (pmol/min/mg)
Production of AFB ₁ -8,9-epoxide:			
Rainbow trout	13°C	81.2 ± 12.8	35.9 ± 4.1
Rainbow trout	28°C	109.3 ± 15.0	66.6 ± 7.1
Zebrafish (female)	28°C	79.0 ± 16.4	11.7 ± 1.4
Production of AFL:			
Zebrafish (female)	28°C	11.2 ± 1.3	9.1 ± 0.3

 $^{^{\}circ}$ n=10 for trout, and n=20 for zebrafish; all assays were run in duplicate, with nonenzymatic background activity subtracted; errors are given as asymptotic standard error. Only female zebrafish were used for this comparison against juvenile rainbow trout, which do not show sexual differences in AFB₁ metabolism prior to maturity.

Discussion

The results demonstrate that zebrafish have the capacity for both Phase I and Phase II metabolism of AFB₁. The major in vivo metabolites excreted into water were AFL and AFL-g, followed by unreacted AFB₁. This metabolic profile is similar to that seen in other fish species, such as the Japanese medaka, rainbow trout, and channel catfish (Toledo et al., 1987; Gallagher and Eaton, 1995; Loveland et al., 1984). AFL, produced by a cytosolic reductase reaction, was recovered in the water at 5 minutes, the earliest time point examined. This has also been observed in the Japanese medaka (Toledo et al., 1987). It is important to note that while AFL is a polar metabolite of AFB₁, it has been shown to be 70% as mutagenic and just as carcinogenic as AFB₁ in rainbow trout, and is also a potent carcinogen in Fischer 344 rats (Coulombe et al., 1982; Schoenhard et al., 1981; Nixon et al., 1981; Bailey et al., 1994). The major conjugation (Phase II) reaction was glucuronidation, which is an important reaction in eliminating hydroxylated compounds in fish (Clark et al., 1991). No evidence of glutathione conjugation was present, as shown by co-chromatography with a glutathione conjugate standard. Formation and excretion of the aflatoxin metabolites occurred rapidly in the zebrafish after i.p. injection.

The *in vitro* assay assessing AFB₁ metabolism again demonstrated the proficiency of AFL formation in zebrafish. This assay also evaluated the ability for AFB₁ epoxidation to the reactive intermediate by trapping the epoxide as the aflatoxinglutathione adduct formed. The results from this assay revealed that zebrafish possess the enzymes necessary for bioactivation of this carcinogen. When compared to trout liver homogenates assayed at 13°C, zebrafish appear to have a comparable Km and only

a 3-fold lower Vmax, which suggests that zebrafish should be a fairly sensitive species for AFB₁ carcinogenesis. Buchmann *et al.* (1993) conducted immunohistochemical analysis to detect the presence of the cytochrome P450 isozyme LMC2 (CYP2K1) in zebrafish. This isozyme is believed to be primarily responsible for the bioactivation of AFB₁ to the epoxide in rainbow trout (Williams and Buhler, 1983). The immunohistochemical analysis of zebrafish using antibodies directed against trout CYP2K1 showed that this isozyme was constituitively expressed in the liver, kidney, skin, and oral mucosa of zebrafish, with lower expression in gills, pseudobranch, intestines, and ovaries. It is possible that this or a similar P450 may be responsible for the observed bioactivation of AFB₁ in zebrafish.

A significant difference in DNA adduct formation between male and female zebrafish was observed in all DNA binding experiments performed. An *in vitro* AFB₁ metabolism assay also confirmed the difference between males and females in their ability to bioactivate AFB₁ to the *exo*-AFB₁ epoxide, which is trapped with high selectivity by the mouse cytosolic glutathione transferase in this assay (Rany *et al.*, 1992b). The *in vitro* assay did not measure differences in the abilities of males and females to produce the *endo*-epoxide, which is not efficiently trapped by the mouse enzyme. It might be that there is a different ratio of *endo*- and *exo*-epoxide production in males and females. Differences between sexes in their ability to metabolize xenobiotics are not unusual. A difference in xenobiotic biotransformation between male and female zebrafish was previously noted by Buchmann *et al.* (1993), where males tended to have higher 7-ethoxyresorufin-O-deethylase activity. This activity is usually associated with the presence of cytochrome P4501A1. The *in vivo* dose-response of

hepatic aflatoxin-DNA adduct formation was linear from 50-400 μ g AFB₁/kg body weight, which is consistent with experiments examining hepatic DNA binding in other fish species following AFB₁ administration (Dashwood *et al.*, 1988; Toledo *et al.*, 1987; and Witham *et al.*, 1982). Using the highest dose, the time-course of aflatoxin-DNA binding showed no statistical difference in binding levels from 1 to 21 days, suggesting slow DNA repair in zebrafish liver. A useful method for comparing DNA binding between species is the CBI (cumulative binding index, units of μ mol chemical bound/mol DNA/mmol chemical administered/kg body weight). A composite CBI calculated from the three DNA-binding experiments is approximately 70,600 in female zebrafish and 35,000 for males. For comparison, the CBI for rat and Japanese medaka is approximately 10,000 and 13,000, respectively, and is 240,000 for trout (Toledo *et al.*, 1987, and references therein). From these data, zebrafish exhibit approximately a 4-fold lower capacity for DNA adduct formation compared to trout, and a 5-fold higher capacity than rat and medaka.

The results from both the investigation of *in vivo* DNA binding after i.p. administration of [³H]AFB₁ and the *in vitro* AFB₁ metabolism assay suggest that zebrafish should be a fairly sensitive model for AFB₁ carcinogenesis, if indeed the amount of DNA adducts present in this species truly correlates with tumor incidence as demonstrated in the trout model (Dashwood, *et al.*, 1992). Not only are zebrafish capable of bioactivating AFB₁ to its active epoxide form, but the DNA adducts that result seem to be persistent over time. However, a study conducted in our laboratory investigating dietary exposure to AFB₁ is providing evidence to the contrary. Zebrafish appear to be quite resistant to the carcinogenic effects of AFB₁ when administered by

the dietary route (Tsai, 1996). A preliminary investigation of DNA binding after dietary administration of [³H]AFB₁ showed only a low level of adduct formation (unpublished results). This dramatic difference is suggestive of a difference in absorption. It appears that if one could get the carcinogen to the target organ, zebrafish might prove quite sensitive. Studies assessing carcinogenicity of i.p. administered AFB₁ have yet to be conducted. It may also be that, while there is evidence for persistence of DNA adduct formation after i.p. administration, these adducts simply do not go on to form initiated cells with any great efficiency in this species. Finally, even if initiation occurs, there could be a lack of promotion/progression of these cells in zebrafish liver. Further studies will be required to test these hypotheses.

Other studies conducted in our laboratory with dietary administration of high amounts of diethylnitrosamine, dimethylnitrosamine, or dimethylbenz[a]anthracene for 6 months have shown limited or no tumor response as well (Tsai, 1996; unpublished results). However, high doses of dietary methylazoxymethanol acetate did manage to elicit almost a 30% tumor response (Tsai, 1996). Detailed metabolism studies have not been conducted after administration of these various carcinogens, so it is uncertain if these low tumor responses are the product of low capacity for bioactivation, high capacity for detoxification and elimination, inability of these compounds when administered in the diet to reach the respective target organs, or some other factor governing the carcinogenic process. Interestingly, zebrafish exposed as embryos or fry to many of these carcinogens are proving to be more sensitive, indicating that zebrafish may be more susceptible to carcinogens at an early stage of development.

In conclusion, adult zebrafish can rapidly metabolize and excrete the potent hepatocarcinogen AFB₁ after intraperitoneal administration. Both *in vitro* and *in vivo* studies verify that this species has the capacity to bioactivate AFB₁ to its reactive intermediate. The DNA adducts that result are suggestive of sensitivity to this carcinogen. These results are in apparent contradiction to the resistance of zebrafish to dietary administration of AFB₁, and suggest a mechanism related to factors other than inherent ability to metabolize and bioactivate this carcinogen.

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CHAPTER 3

CYP1A INDUCTION BY β -NAPHTHOFLAVONE, AROCLOR 1254, AND 2,3,7,8-TETRACHLORODIBENZO-p-DIOXIN, AND ITS INFLUENCE ON AFLATOXIN B₁ METABOLISM AND DNA BINDING IN ZEBRAFISH (Danio rerio)

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Abstract

This study investigated the response of cytochrome P450 1A (CYP1A) in the zebrafish (Danio rerio) following exposure to Aroclor 1254, β -naphthoflavone (β NF), and 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD), and then investigated TCDD modulation of aflatoxin B₁ (AFB₁) metabolism and hepatic AFB₁-DNA adduction. Aroclor 1254 fed at 500 ppm for 1 to 9 days or intraperitoneal (i.p.) injection of 75 -200 mg Aroclor 1254/kg body weight failed to induce CYP1A protein or associated 7ethoxyresorufin-O-deethylase (EROD) activity. Dietary β NF at 500 ppm for 3 or 7 days induced CYP1A protein and EROD activity approximately 3-fold above controls. A single i.p. injection of 150 mg β NF showed maximal induction of CYP1A protein and EROD activity by 24 hours, both of which decreased rapidly during the next 6 days. A more refined study showed maximum CYP1A protein levels between 24 and 36 hours, while EROD activity had an observable peak at 16 hours. CYP1A and EROD activity showed dose-responsiveness following single i.p. administration of 25, 50, 100, or 150 mg β NF/kg body weight. Dietary exposure to 0.75 ppm TCDD for 3 days also significantly induced CYP1A. The effect of TCDD on the metabolism of [3H]AFB, in zebrafish was then investigated. The major [3H]AFB, metabolites excreted in water over 24 hours in the control group were aflatoxicol, aflatoxicol-glucuronide, and parent AFB₁. By contrast, the predominant metabolites in the TCDD-pretreated group were aflatoxicol-M₁-glucuronide, aflatoxicol, aflatoxin M₁ plus aflatoxicol-M₁ unresolved, aflatoxicol-glucuronide, and parent AFB₁. Surprisingly, hepatic AFB₁-DNA adduction was approximately 4-fold higher in the TCDD treated group compared to controls. This significant difference could not be explained by increased capacity for bioactivation of AFB₁ as measured by an *in vitro* AFB₁ metabolism assay. However, it was demonstrated that zebrafish have the capacity to bioactivate aflatoxin M_1 to a reactive intermediate, and it is proposed that secondary bioactivation of this genotoxic intermediate may be responsible for the increased DNA binding.

Introduction

Fish are becoming important models in many areas in science, including developmental biology, genetics, neurobiology, aquatic toxicology, and carcinogenesis (Powers, 1989). The trout model has been and continues to be extensively characterized as an alternative nonmammalian vertebrate model in carcinogenesis research (for review see Bailey et al., 1996). The use of smaller fish as experimental models is also assuming prominence because they have several advantages over the trout model such as their small size, ability to spawn regularly, reduced husbandry costs, and their overall hardiness (Hawkins et al., 1988). Many small fish species, such as the guppy, Japanese medaka, and zebrafish are proving to be susceptible to numerous carcinogens, including diethyl- and dimethylnitrosamine, 2-acetylaminofluorene, nitrosomorphiline, and methylazoxymethanol acetate (Sato et al., 1973; Khudoley, 1984; Fournie et al., 1987; Nakazawa et al., 1985; Hawkins et al., 1986; Stanton, 1965). Zebrafish are a particularly attractive fish model because they are also being extensively developed in both genetics and developmental biology. Studies in our laboratory investigating the sensitivity of zebrafish to carcinogens are demonstrating that this species is a complex model. Adult zebrafish appear to be somewhat resistant to carcinogens administered in

the diet, but zebrafish are susceptible to carcinogen exposures in the water, particularly when exposed as embryos and fry (Tsai, 1996; unpublished results).

Induction of cytochrome P450 1A (CYP1A) in fish is being investigated as a biomarker of exposure to certain compounds commonly found in the environment such as polyaromatic hydrocarbons (PAHs), polychlorinated biphenyls (PCBs), and halogenated dibenzo-p-dioxins and dibenzofurans (for review see Goksoyr, 1995). While fish appear to lack the equivalent of inducible phenobarbital-type P450s, most fish species do possess a CYP1A-like P450 that can be effectively induced to a high level (Goksoyr et al., 1991; Stegeman, 1989; Stegeman and Hahn, 1994). CYP1A is not normally constituitively expressed in significant levels. Following induction, it is primarily located in hepatic tissue, although in fish, it can also be found in extrahepatic tissues such as gill, kidneys, gut and heart (Buchmann et al., 1993; Stegeman et al., 1989; Husoy et al., 1994). Induction of CYP1A occurs following exposure to a wide range of compounds, including the classic inducer 3-methylcholanthrene, many flavones and indoles, and the PAHs and PCBs. In mammals, the Ah-receptor is the mediator of induction, and is known to be responsible not only for the induction of CYP1A1, but a host of other enzymes including quinone-oxido-reductase, glucuronsyltransferase, glutathione-S-transferase, CYP1A2, and CYP2B1 (Poland, 1979; Hankinson, 1995).

The enhanced metabolic capabilities following exposure to one of the many CYP1A-inducing agents may result in altered metabolism of other xenobiotics including procarcinogens. One of the most extensively characterized procarcinogens is the potent mycotoxin aflatoxin B₁ (AFB₁), which requires bioactivation to the reactive intermediate, the AFB₁-8-9-epoxide. While both the *endo-* and *exo-*AFB₁ epoxide may be produced,

it is primarily the exo-epoxide stereoisomer that is responsible for binding to DNA (Raney et al., 1992a; 1992b; Eaton and Gallagher, 1994). This mycotoxin is carcinogenic in many species, including rat, human, and rainbow trout, the most sensitive species. Numerous investigations have explored the modulation of aflatoxin B₁ metabolism and carcinogenesis by compounds known to act via the Ah receptor. Studies with rainbow trout have shown that pre-exposure to the synthetic flavone β naphthoflavone (β NF), the PCB mixture Aroclor 1254, and the natural plant constituent indole-3-carbinol reduces AFB, carcinogenesis by altering the metabolism of AFB, and reducing DNA adduction (Shelton et al., 1986; Goeger et al., 1986; Nixon et al., 1984; Takahashi et al., 1995; 1996). An in vitro investigation of β NF-induced channel catfish microsomes showed induction of CYP1A enhanced detoxification of the carcinogen, but did not affect bioactivation of AFB, to its reactive intermediate (Gallagher and Eaton, 1995). Other studies investigating β NF-induced rabbit pulmonary and hepatic microsomes and β NF-induced guinea pig liver, kidney, and lung microsomes have also shown increased formation of the less toxic metabolite aflatoxin M, and reduced DNA adduction (Daniels and Massey, 1992; Liu, et al., 1993). Although 2,3,7,8tetrachlorodibenzo-p-dioxin (TCDD) is the most potent Ah receptor agonist known, surprisingly few studies have investigated the modulation of AFB, metabolism and DNA adduction by pre-exposure to TCDD. The purpose of this study was to characterize CYP1A induction in zebrafish by Aroclor 1254, β NF, and TCDD. After characterizing CYP1A induction, further studies were conducted to investigate the in vivo modulation of AFB₁ metabolism and hepatic DNA adduction following dietary exposure to TCDD at a concentration known to exert biological effects on metabolizing enzymes.

Materials and Methods

Chemicals

AFB₁ and AFM₁ were purchased from Sigma Chemical Company (St. Louis, MO), and assessed for purity by thin layer chromatography (TLC); [³H]AFB₁ was from Moravek Biochemicals (Brea, CA) and the chemical and radiopurity was checked by TLC followed by radioscanning; βNF was from Fluka Chemical Corp. (Ronkonkoma, NY); Aroclor 1254 was from Monsanto Company (St. Louis, MO); TCDD was from ANALABS (New Haven, CT); 7-ethoxyresorufin and resorufin were from Molecular Probes, Inc. (Eugene, OR); RNase, DNase free were from Boehringer Mannheim Biochemicals (Indianapolis, IN); HPLC grade acetonitrile, methanol, tetrahydrofuran, and J.T. Baker C₁₈ Empore extraction disks were from VWR (Seattle, WA); Hoechst #33258 was from Calbiochem-Behring Corp. (La Jolla, CA); aflatoxin HPLC standards aflatoxicol (AFL), aflatoxicol-glucuronide (AFL-g), aflatoxin M₁ (AFM₁), and aflatoxicol-M₁ (AFL-M₁) were prepared in our laboratory by previously published methods (Loveland *et. al.*, 1983; 1984).

Animal care and maintenance

Adult (sexually mature) zebrafish were reared and maintained in the Food Toxicology and Nutrition Laboratory in aerated 29 gallon aquariums with a controlled temperature of 26°C (± 1) and a 14 hour light: 10 hour dark photoperiod. During the week, fish were fed twice daily with TetraMin Staple Food (flakes) each morning, and

a combination of Oregon Test Diet (OTD) (Sinnhuber et al., 1977) and brine shrimp in the afternoon on Monday, Wednesday, and Friday, and TetraMin Staple Food and brine shrimp on Tuesday and Thursday afternoons. On weekends, fish received only one feeding per day consisting of OTD and brine shrimp. For dietary exposures, fish were fed one feeding of OTD or the test reagent in OTD per day during the course of the treatment. For the Aroclor 1254 dietary exposure only, fish were fed a control diet consisting of the purified casein diet (PC diet) or control diet containing Aroclor 1254 (DeKoven et al., 1992).

Preparation of liver tissue

All fish were fasted for 24 hours before receiving the appropriate treatment. When fish were sampled, livers were immediately excised from the zebrafish, the gall bladders carefully removed, and the unrinsed livers placed on ice until the appropriately sized pools of tissue were obtained. The samples were then flash frozen in liquid nitrogen and stored at -80°C. Before use, livers were homogenized in ice-cold buffer containing 0.1 M potassium phosphate (pH 7.25), 20% glycerol, 1 mM ethylenediaminetetraacetic acid, 1 mM dithiolthreitol, 0.1 mM butylated hydroxytoluene, and 0.1 mM phenylmethylsulfonylfluoride. The samples were centrifuged for 10 minutes at 1,000x g and the supernatant decanted for protein work. When DNA binding was also being investigated, the pellet was saved for DNA isolation and purification.

Investigation of CYP1A induction by Aroclor 1254, \(\beta NF, \) and TCDD

A total of 18 male and 18 female adult zebrafish were fed 500 ppm Aroclor 1254 twice per day and 3 males and 3 females each were sampled on 1, 2, 3, 5, 7, or 9 days. Four males and 4 females were fed the control PC diet twice per day and 1 male and 1 female sampled at days 2, 3, 5 and 7. When dietary exposure failed to show induction of CYP1A, 6 adult, female fish were i.p. injected with 100 or 200 mg Aroclor 1254/kg body weight, 4 were injected with carrier dimethylsulfoxide (DMSO), and all were sampled 24 hours later.

To confirm results from a preliminary investigation which showed CYP1A induction after i.p. exposure to βNF , a time-course study was undertaken. Thirty adult female zebrafish were i.p. injected with 150 mg β NF/kg body weight and sampled at day 1, 2, 3, 4, and 7; 6 controls were injected with DMSO and sampled at day 1, 2, and 3. A concurrent dietary exposure to 500 ppm of β NF was also conducted to investigate whether dietary exposure to this flavone was capable of inducing CYP1A in zebrafish. Sixteen adult female fish were fed 500 ppm β NF and 8 fish fed control diet for 3 or 7 days. To further pinpoint the time of maximum CYP1A induction by β NF, 32 adult female zebrafish were i.p. injected with 150 mg/kg body weight of β NF in DMSO and sampled at 8, 16, 24, or 36 hours after injection. Twenty-four female zebrafish were also injected with 25, 50, or 100 mg/kg body weight of β NF in DMSO and sampled 24 hours later to try to establish a dose response (the 24 hour group in the time-course served as the 150 mg β NF/kg body weight dose). To evaluate if the use of DMSO instead of corn oil as a vehicle influenced the kinetics of induction, 40 females were i.p. injected with 150 mg/kg body weight of β NF in corn oil and sampled 24, 36, 48, 72, and 96 hours after injection. Lastly, to verify previous results demonstrating a lack of detectable CYP1A following exposure to Aroclor 1254, 16 female fish were i.p. injected with 75 or 150 mg Aroclor 1254/kg body weight using DMSO as a carrier. Ten females were i.p. injected with DMSO and 8 females were injected with corn oil and sampled 24 hours later to serve as controls for these experiments.

Finally, to verify that TCDD induces CYP1A in the zebrafish, 16 adult female zebrafish were fed 0.75 ppm TCDD or control diet for 3 days and sampled on the fourth day. This dosage was based on a previous study by Buchmann *et al.* (1993) which also investigated CYP1A in zebrafish.

TCDD modulation of aflatoxin DNA-binding

Sixty-four adult female zebrafish were fasted for 24 hours and then fed control OTD or OTD diet containing 0.75 ppm TCDD for 3 days. The day after feeding was completed, 8 of the control fish and 8 of the TCDD-treated fish were sampled for representative assessment of CYP1A levels and EROD activity at the time of AFB₁ dosage. Then, 12 of the control and 12 of the TCDD-treated fish were i.p. injected with DMSO and the other 12 control and 12 TCDD-treated fish were i.p. injected 400 µg/kg body weight of [³H]AFB₁ (16.6 Ci/mmol) in DMSO. Fish were rinsed with 1 ml of water to account for residual skin contamination before placement in respective 5 gallon buckets containing approximately 16 liters of aerated water. Rinse water accounted for less than 4% of the injected dosage. Livers were sampled 24 hours after injection.

DNA was isolated and purified using a modification of Strauss (1991) (modification found in Troxel et al., Chapter 2 this thesis). DNA was quantified using the microfluorometric procedure of Cesarone et al. (1979) with a Hoefer TKO100 DNA fluorometer. The amount of [³H]AFB₁ bound to DNA was determined after hydrolyzing the DNA by heating the samples at 70°C with equal volumes of 1 M perchloric acid for 20 minutes and then counting with a Beckman LS 7500 scintillation counter.

TCDD modulation of AFB₁ metabolism

Twelve adult female zebrafish were fasted for 24 hours and then fed control OTD or OTD diet containing 0.75 ppm TCDD for 3 days. The day after feeding was completed, 3 control and 3 TCDD-treated fish were i.p. injected with DMSO, whereas the remaining 3 control and 3 TCDD-treated fish were i.p. injected with 400 µg/kg body weight of [³H]AFB₁ (16.6 Ci/mmol) in DMSO. Fish were rinsed with 1 ml of water to remove any residual radioactivity before placement in individual beakers containing 50 ml of water. Rinse water contained less than 5% of the administered dose except for fish 2 of the AFB₁/TCDD treatment group, which contained approximately 8% of the injected dosage. At various time points, fish were removed to fresh beakers of water, the previous water samples were collected, and an aliquot was taken for scintillation counting. At 24 hours, the livers of the fish were sampled and treated as described earlier for enzyme analysis and quantification of DNA adduction.

Water samples were extracted with C_{18} extraction disks and metabolite analysis performed by HPLC as described (Troxel *et al.*, Chapter 2 this thesis). The amount of radioactivity remaining after filtration by the C_{18} filter (i.e., the amount not trapped by

the filter) was higher than expected in some samples, so these water samples were refiltered with another C₁₈ filter. Some additional radioactivity was trapped in the 7 and 24 hour samples of the AFB₁/TCDD treated group, suggesting incomplete adsorption in the first filtration. However, there still was a high amount of radioactivity remaining unbound in these doubly filtered water samples. To investigate the nature of this material, the 7 and 24 hour water samples from both AFB₁-treated groups were treated with 0.2 M sodium acetate buffer (pH 5.0), buffer containing β -glucuronidase (4000) units/ml), or sulfatase (20 units/ml, with 40 mM D-saccharic acid-1,4-lactone added to inhibit β -glucuronidase activity) (Fong et al., 1993). The samples were incubated at 37°C for 18 hours before extracting twice with 2 volumes of ethyl acetate. When analysis of the results showed some glucuronidation in the 7 and 24 hour sample from the AFB,/TCDD treated group and the 24 hour sample from the AFB, group, up to 50 ml of the water samples were evaporated down to 1 ml, and the samples were then injected into the HPLC for further analysis. Under these HPLC conditions, AFM, and AFL-M₁ were not fully resolved, and therefore are reported together in the analysis.

Immunoblotting

Using bovine serum albumin as the standard, total protein was quantified by the method of Lowry et al. (1951). Proteins were separated by SDS-PAGE (Laemmli, 1970) and electrophoretically transferred to nitrocellulose membranes (Towbin et al., 1979). Due to the large number of samples to compare, after verification that only one cross-reacting band was present in blots analyzed for CYP1A protein, subsequent analysis of CYP1A was limited to dot blots. The blots were incubated with rabbit anti-

trout IgG against LM_{4b} (CYP1A) at a concentration of 2 μ g/ml. The membranes were next probed with goat anti-rabbit secondary antibody conjugated to horseradish peroxidase. The P450 isozyme was detected using the ECL chemiluminescence detection kit. Densitometry analysis was performed with an HP ScanJet flatbed scanner with NIH Image version 1.54 software (Wayne Rasband, NIH, public domain). Previously quantified β NF-induced trout liver microsomes were used as standards on the blots, and all blots were normalized with the same sample of β NF-induced zebrafish liver homogenate.

Enzyme assays

EROD activity was determined by the method of Prough *et al.* (1978). The assay was conducted at 30°C with a substrate concentration of 2 μ M.

The *in vitro* AFB₁ metabolism assay was conducted using the method of Monroe and Eaton (1987), with modifications by Takahashi *et al.* (1996). This assay can indirectly measure bioactivation of AFB₁ by trapping the reactive intermediate with glutathione. Previous studies in which trout microsomes were incubated with AFM₁ instead of AFB₁ have shown the formation of another glutathione adduct with a different HPLC retention time than that of the AFB₁-glutathione adduct, suggesting that trout also bioactivate AFM₁ to a reactive intermediate (data not shown). In this experiment, the metabolism of both AFB₁ and AFM₁ was investigated in zebrafish liver homogenates. The final assay concentration of AFM₁ or AFB₁ was 80 μ M, and 0.51 mg/ml of supernatant protein from liver homogenates was used. Assay conditions consisted of a 2 minute pre-incubation at room temperature before the addition of the glutathione and

NADPH. Incubations were carried out at 28°C for 45 minutes before termination with ice-cold 2 M acetic acid and internal standard AFG₁. Preliminary studies showed that this time point was in the linear range of time.

Statistical analysis

Statistical analysis was performed with SAS, version 6.10 (SAS Institute Inc., 1989). Differences in the means between two groups were determined using the t-test procedure (equal or unequal variance t-test). Time or dose response data were analyzed using analysis of variance (GLM procedure), followed by polynomial trend analysis. The dietary β NF study was analyzed by two way analysis of variance (days and β NF dose effects). A p value less than 0.05 was considered significant in all analyses.

Results

Investigation of CYP1A induction in adult zebrafish by Aroclor 1254, \(\beta NF \), and TCDD

The dietary exposure of adult zebrafish to 500 ppm Aroclor 1254 for up to 9 days failed to increase CYP1A protein quantities above control levels, and did not measurably induce hepatic EROD activity. A subsequent study investigating the induction of CYP1A following i.p. injection of 100 or 200 mg Aroclor 1254/kg body weight also showed no response to this common inducer. A second attempt to induce CYP1A following i.p. injection of 75 or 150 mg Aroclor 1254 was again unsuccessful.

Thus Aroclor 1254 failed to evoke a detectable response in the zebrafish by either dietary or i.p. treatment.

By contrast, a time-course investigation of CYP1A induction following i.p. injection of 150 mg β NF/kg body weight showed strong and maximal induction of both CYP1A protein and EROD activity by 24 hours, the first time-point examined following injection, with a curvilinear decrease over time (linear, p=0.0001; quadratic, p<0.004) (Figure 3.1, panel A). There was a rapid decrease in both protein and activity during the remaining course of the week, with barely detectable levels of EROD activity and non-detectable levels of protein at day 7. A dietary exposure of 500 ppm β NF for 3 or 7 days demonstrated an approximate 3-fold increase in protein levels and EROD activity above controls (p<0.04, β NF main effect), but there was no statistical difference between the days (p=0.7) (Figure 3.1, panel B). Noteworthy is that low but detectable levels of CYP1A were observed in the controls from the dietary exposure, whereas control values in the injection experiment were non-detectable. This is suggestive of a dietary component possessing some CYP1A-inducing activity.

Additional studies were conducted to further investigate the response of zebrafish CYP1A following exposure to β NF. Another time-course experiment using a narrower range of time was performed, because CYP1A was already maximally induced at the first time point examined in the previous time-course experiment. This study showed that protein levels displayed an increasing linear response over time (p=0.003), and were highest between 24 and 36 hours following injection, while the EROD activity exhibited a significant curved response (quadratic, p=0.035), with an observed peak at 16 hours (Figure 3.2, panel A). It is interesting to note that the amount of induction

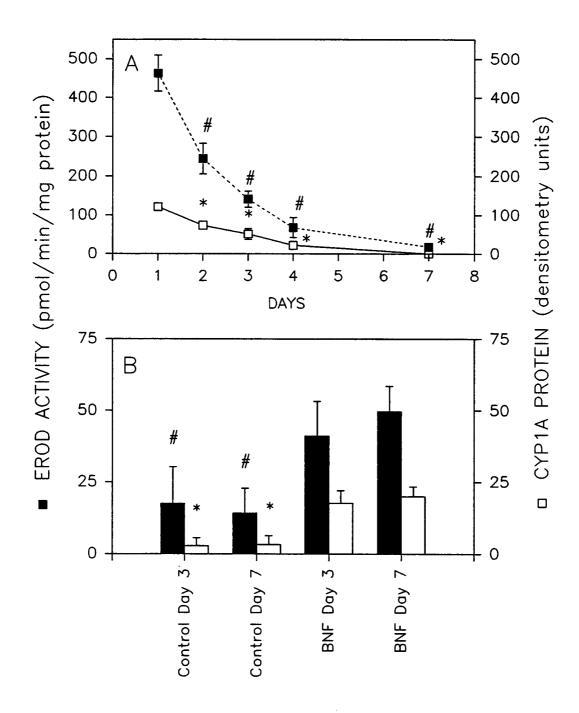


Figure 3.1. Induction of CYP1A protein and EROD activity in adult female zebrafish following i.p. or dietary administration of β NF. A) Single i.p. injection of 150 mg β NF/kg body weight (data are means \pm SEM from 3 pools of 2 fish each) B) dietary exposure to 500 ppm β NF (data are means \pm SEM from 4 pools of 2 fish each for treated, and 2 pools of 2 fish each for controls). * and # designate statistical difference (p<0.05) from the observed peak for protein induction (*) and EROD levels (#).

Figure 3.2. Induction of CYP1A protein and EROD activity in adult female zebrafish by β NF following different treatments. A) single i.p. administration of 150 mg β NF/kg body weight with sampling up to 36 hours after injection B) i.p. administration of 25, 50 100, or 150 mg β NF/kg body weight and sampled 24 hours later C) single i.p. administration of 150 mg/kg body weight of β NF in corn oil instead of the carrier DMSO, sampled up to 4 days following injection. All data are means \pm SEM for 4 pools of 2 fish each). * and # designate statistical difference (p<0.05) from the observed peak for protein induction (*) and EROD levels (#).

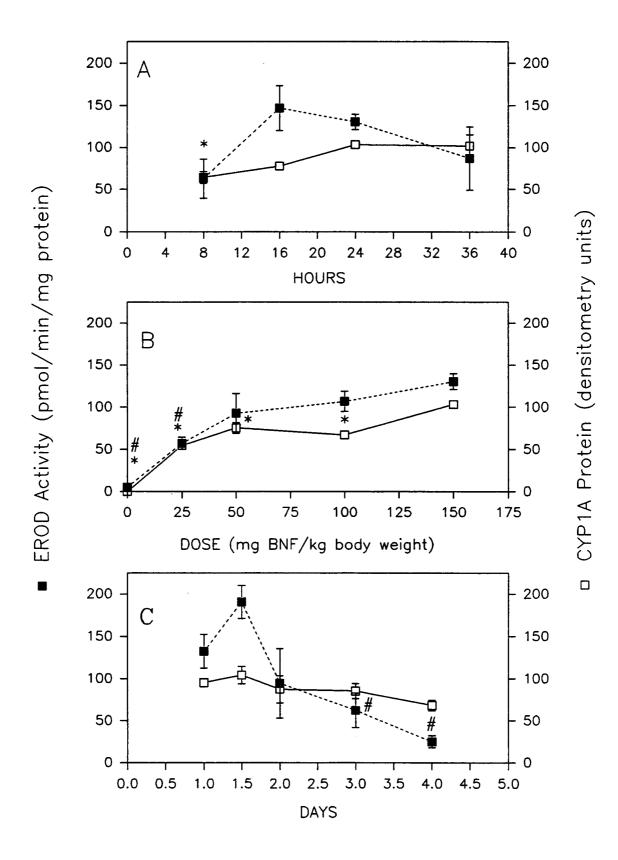


Figure 3.2.

documented in this experiment is considerably less than that observed in the first study. It should also be noted that these samples showed high variability between groups, especially at the 36 hour time-point, which might explain the discrepancy with maximal induction of EROD activity occurring before peak induction of protein. Intraperitoneal injection of 25, 50, 100, or 150 mg β NF/kg body weight provided dose-responsive induction of protein and EROD activity (Figure 3.2, panel B). The response to β NF was also investigated by conducting another time-course experiment following i.p. injection of 150 mg/kg body weight of β NF in corn oil instead of DMSO (Figure 3.2, panel C). Over the range of four days, there was a decreasing trend for EROD activity (p=0.0004), with an observed peak at 36 hours. This time point, however, was not statistically different from the 24 hour time point (p=0.1), which is consistent in the experiments with DMSO as a carrier. As in the experiment in Figure 3.2, panel A, levels of CYP1A protein showed less dramatic changes than EROD activity over the period examined.

TCDD, the most potent Ah receptor agonist yet described, was found to be a potent inducer of hepatic CYP1A in this species as well. Zebrafish fed 3 days with 0.75 ppm TCDD in OTD had an EROD activity approximately 17-fold higher than controls (391 \pm 29 pmol/min/mg protein compared to 23 \pm 5 pmol/min/mg protein). Protein levels were also greatly induced in TCDD-treated fish (96 \pm 17 densitometry units versus non-detectable levels in controls).

TCDD modulation of AFB, metabolism in vivo

Early excretion kinetics of AFB₁ metabolites following i.p. administration of $[^3H]AFB_1$ was assessed by measuring the amount of radioactivity recovered in the water over 24 hours. The amount of radioactivity did not significantly differ between treatment groups. Approximately $52 \pm 13\%$ of the administered radioactivity was recovered by 24 hours in the TCDD/AFB₁ treatment group, while $45 \pm 7\%$ was accounted for in the AFB₁ treatment group (Figure 3.3). These data show that, overall, early excretion kinetics of $[^3H]AFB_1$ were not strongly affected by preexposure of zebrafish to TCDD.

However, whereas the total amount of radioactivity recovered in the water was similar, the actual metabolic profile of AFB₁ was different between the two groups. The AFB₁ treatment group exhibited a similar metabolic profile to that seen in the earlier metabolism study (Troxel *et al.*, Chapter 2 in this thesis). Aflatoxicol (AFL) accounted for approximately 17% of the original dose administered, followed by unreacted AFB₁ at 6.1% and aflatoxicol-glucuronide (AFL-g) at 5.3% (Figure 3.4, panel A). Less than 1% of the administered dose was recovered as aflatoxin M₁ (AFM₁), aflatoxicol M₁ (AFL-M₁), or aflatoxicol-M₁-glucuronide (AFL-M₁-g) combined. As was expected in the TCDD/AFB₁ treatment group, there was a higher production of AFM₁/AFL-M₁ and of the glucuronides. AFL-M₁-glucuronide was the predominate metabolite, accounting for 8.6% of the original dose, followed by AFB₁ and AFL with approximately 7.5% each, AFM₁ and AFL-M₁ with 5.6%, and lastly AFL-g with 4.0% (Figure 3.4, panel B). Treatment of the water samples with sulfatase demonstrated that

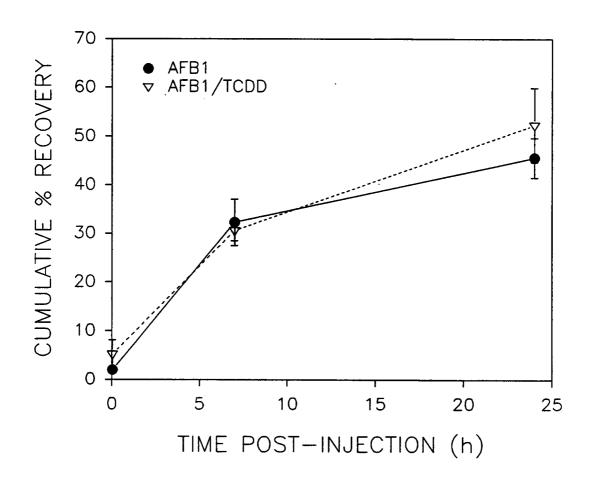


Figure 3.3. The cumulative percentage of radioactivity recovered in the water within 24 hours in control and TCDD-treated zebrafish following i.p. administration of [³H]AFB₁.

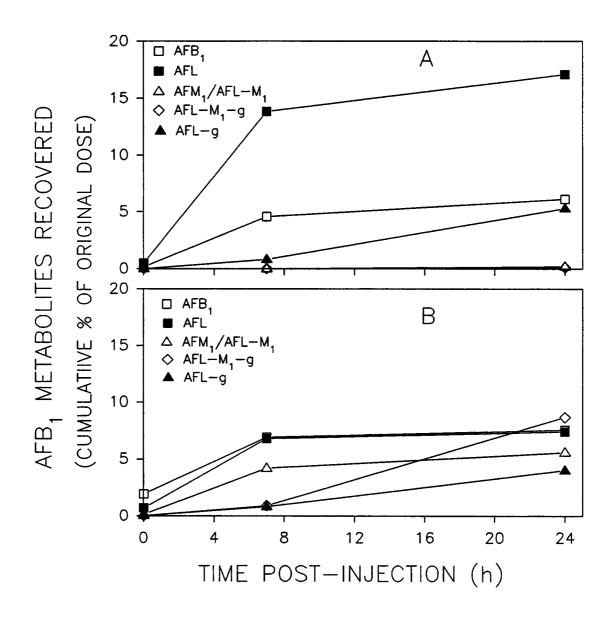


Figure 3.4. The cumulative percentage of individual metabolites recovered in water over 24 hours following i.p. administration of [³H]AFB₁, expressed as a percentage of the original dose administered A) in the control-fed group B) in the TCDD-fed group.

sulfate conjugation represented only a minor metabolite (at most representing less that 0.5% of the administered dose) in the zebrafish.

Figure 3.5 shows the percentage of each metabolite recovered at each time point as a percentage of the original dose administered. Overall, the total percentage of recovery of the metabolites varied slightly between the groups, with 29% of the dose recovered in the AFB₁ group, and 33.2% in the TCDD/AFB₁ group. The slight increase in metabolites recovered in the TCDD/AFB₁ group occurred primarily at the 24 hour time point, where there was the increased production and excretion of the glucuronides of AFL and AFL-M₁. The greater production of AFL-M₁-g compared to AFL-g might reflect substrate availability, or perhaps the more polar AFL-M₁ may be preferentially glucuronidated over AFL.

TCDD modulation of in vivo hepatic AFB-DNA adduction and in vitro activation of AFB₁ and AFM₁

The *in vivo* hepatic AFB-DNA adduction at 24 hours after i.p. injection of 400 μg [3 H]AFB₁/kg body weight was determined to be almost 4-fold higher in the group pre-exposed to TCDD than in the control group (1660 \pm 175 pmol AFB₁/mg DNA versus 438 \pm 109 pmol AFB₁/mg DNA, p < 0.05) (Table 3.1). To further investigate the cause of this notable difference in adduction levels, *in vitro* AFB₁ and AFM₁ metabolism studies were conducted (Table 3.1). The results from these metabolism experiments showed no statistical difference between the control or TCDD-treated groups in their ability to bioactivate AFB₁ or AFM₁ to the respective reactive intermediate. There was also no statistical difference between the two groups in their

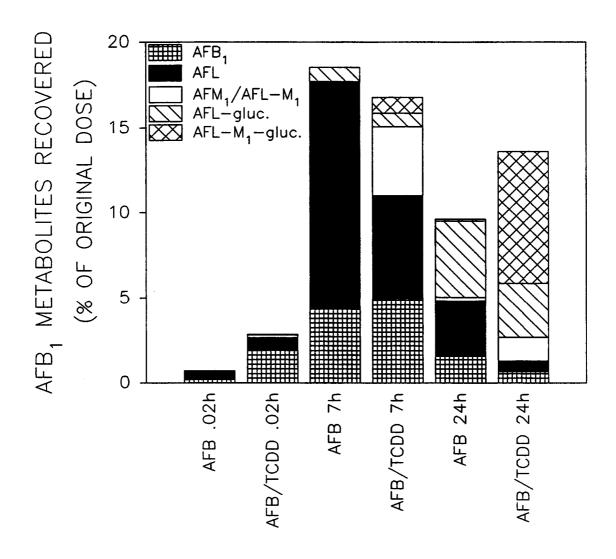


Figure 3.5. The percentage of all metabolites recovered in water at each time point, expressed as a percentage of the original dose in the control-fed group and in the TCDD-fed group.

Table 3.1. Effects of TCDD pretreatment on *in vivo* hepatic [³H]AFB₁-DNA adduction and *in vitro* AFB₁ and AFM₁ bioactivation and metabolism^a.

	AFB ₁	TCDD/AFB ₁
*hepatic DNA adduction		
pmol AFB ₁ /mg DNA	438.2 ± 109.1	1660.2 ± 175.4
baflatoxin-8,9-epoxide		
pmol/min/mg protein	13.5 ± 5.4	9.5 ± 1.0
aflatoxin M ₁ (AFM ₁)		
pmol/min/mg protein	1.0 ± 0.3	24.2 ± 3.0
aflatoxicol (AFL)		
pmol/min/mg protein	9.2 ± 2.4	8.3 ± 0.8
^c aflatoxin M ₁ -8,9-epoxide		
pmol/min/mg protein	8.2 ± 3.1	6.2 ± 0.6

^{*} data are means ± SEM from 3 pools of 4 fish each, and 1 pool of 3 fish each

^b assays were run in duplicate with a final substrate concentration of 80 μ mol AFB₁, with nonenzymatic background subtracted

[°] assays were run singly, with a final substrate concentration of 80 μ M AFM₁, with nonenzymatic background subtracted. The epoxide was trapped as the glutathione adduct similarly to the AFB₁-8,9-epoxide.

capability of metabolizing AFB_1 to the primary metabolite AFL. However, the TCDD-treated group did have a 22-fold increase (p=0.0001) in their capability to metabolize AFB_1 to AFM_1 , a reaction believed to be mediated by CYP1A in zebrafish as has been shown for trout (You *et al.*, unpublished results).

Effects of TCDD on CYP1A protein and EROD activity.

CYP1A protein levels and EROD activity were measured at day 0 (the time that the AFB, dosage was administered) in eight control and eight TCDD-treated fish to provide a representative value of CYP1A induction in the zebrafish at the time of AFB, dosing. The results were in accordance with the preliminary TCDD-dietary exposure already described earlier in this section. The average EROD activity was 337 \pm 10 pmol/min/mg protein, and protein levels were 128 ± 4 densitometry units. In this experiment, no CYP1A activity was detectable in the controls. CYP1A induction was also measured in the supernatant from the samples used in the DNA-adduction and metabolism experiments. Protein levels and EROD activity were non-detectable in the samples that received no TCDD treatment (i.e., the groups fed control diet and injected with either DMSO or [3H]AFB₁). CYP1A protein levels and EROD activity were comparable between the TCDD-treated groups (i.e., the groups fed TCDD and injected with DMSO or [3H]AFB₁). The TCDD-treated group that received [3H]AFB₁ had a 1.6fold higher mean EROD activity than the TCDD-treated group injected with carrier only $(550 \pm 64 \text{ pmol/min/mg protein versus } 344 \pm 68 \text{ pmol/min/mg protein})$, but this difference in mean activity did not achieve significance (p=0.07) (Figure 3.6). The

results verify that TCDD treatment in fact did induce hepatic CYP1A in fish used for the AFB₁-DNA binding experiment.

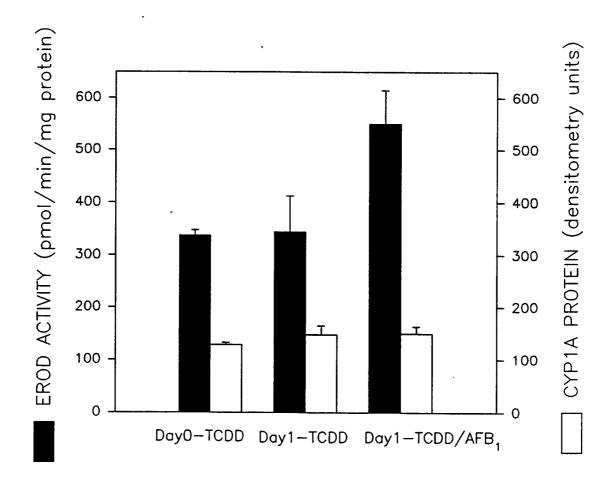


Figure 3.6. Levels of CYP1A protein and associated EROD activity in the supernatant from zebrafish liver homogenates used in the metabolism and DNA adduction experiments. (Data are means \pm SEM from 3 pools of 4 fish each, and 1 pool of 3 fish each).

Discussion

CYP1A in zebrafish

A cytochrome P4501A1-like protein has been found in all fish species so far investigated, including zebrafish. Using anti-trout P4501A1 IgG, Buchmann *et al.* (1993) noted a single cross-reactive band from zebrafish liver microsomes following dietary exposure to TCDD, and Collodi *et al.* (1994) found 2 cross-reactive bands (50,000 and 54,000 kDa) from zebrafish liver homogenates following exposure to a static water bath containing TCDD. Our results showed that a single cross-reacting band was induced in zebrafish liver homogenates following exposure *in vivo* to β NF or TCDD using this same antibody. Interestingly, both an *in vivo* static bath exposure of zebrafish to 50 μ g/l of β NF for 48 hours and zebrafish liver cells exposed to β NF *in vitro* showed no detectable induction of a CYP1A-like protein (Collodi *et al.*, 1994; Miranda *et al.*, 1993). Our studies, however, demonstrated that zebrafish CYP1A was indeed responsive to this flavone in the whole animal following i.p. administration or dietary exposure. The explanation for this difference is unknown.

BNF induction of CYP1A

Induction of the CYP1A-like protein and associated EROD activity following exposure to β NF is similar to that seen in other fish species, including the rainbow trout, Japanese medaka, and goldfish (Zhang et al., 1990; Schell et al., 1987; Gooch and Matsumura, 1983). The decreased levels of activity seen between the first and

second time-course experiments is not too surprising. While both experiments were conducted using fish from the same spawning, the experiments were not performed concurrently. The levels and inducibility of CYP1A in fish can vary greatly depending on the reproductive status of the fish (Förlin and Haux, 1990; Larsen et al., 1992). Courtenay et al. (1994) investigated CYP1A mRNA expression in Atlantic tomcod and found varying degrees of CYP1A mRNA expression in prespawning, spawning, and spent male and female tomcod. The consequences of continuous spawning in such fish as the zebrafish are not known at this time.

Lack of Aroclor 1254 induction of CYP1A

The lack of induction of CYP1A in zebrafish liver following both dietary exposure and i.p. administration of Aroclor 1254 is puzzling. This commercial mixture of PCBs containing 54% chlorine is usually quite effective as a 3-methylcholanthrene-type inducer in most species. Only seldom is non-responsiveness to this mixture documented. One example is the study by Yawetz et al. (1992), in which exposure of a Mediterranean mollusc species to Aroclor 1254 increased overall P450 levels in the digestive gland, but actually decreased EROD activity. In another example, redfish receiving a single i.p. injection of Aroclor 1254 did not show any increase in microsomal mixed function oxidases (Stahl et al., 1984). Studies in fish using high doses of β NF, benzo[a]pyrene, or planar PCBs have demonstrated that high doses of these inducers can actually lead to inhibition of activity or mRNA of the very protein being induced (Gooch et al., 1989; Goddard et al., 1987; Haasch et al., 1993; Melancon and Lech, 1983). In the present study, not only was EROD activity non-

detectable, but protein levels were also not induced above control values, even at the lowest dose (75 mg/kg body weight) investigated. Another possible explanation might be that induction by this PCB mixture could be delayed, and simply was not detected by 24 hours following i.p. administration. The dietary exposure, however, was continued for 9 days with continuous feeding, and was also unsuccessful at inducing CYP1A. The Ah receptor has not yet been isolated in zebrafish. While the assumption is that the mechanism of induction of CYP1A in zebrafish is mediated by the Ah receptor, this has not been verified. If the Ah receptor is indeed present, it might be that the agonist binding site is more selective in this species.

TCDD modulation of AFB, metabolism

The *in vivo* metabolism of AFB₁ in zebrafish observed in this experiment was similar to the metabolic profile documented in a previous study (Troxel *et al.*, Chapter 2 of this thesis). The predominant metabolites excreted into water over 24 hours were again AFL, AFB₁, and AFL-g. The AFB₁ metabolites excreted by the TCDD pretreated group consisted of AFL, AFB₁, and AFL-g, but additionally included AFM₁/AFL-M₁ and the glucuronide of AFL-M₁. It is believed that CYP1A1 is the enzyme responsible for production of AFM₁, and is postulated that AFM₁ can become further hydroxylated to form AFL-M₁, which can than be glucuronidated, as is the case in rainbow trout (Loveland *et al.*, 1983; 1984; Goeger, *et al.*, 1988; You, publication in progress). Therefore it is no surprise to see increased levels of AFM₁/AFL-M₁ and the glucuronide of AFL-M₁.

The *in vitro* metabolism experiments showed increased formation of AFM₁ in the TCDD-treated liver homogenates, confirming the results of the *in vivo* study. Although there was a significant difference in the *in vivo* production of AFL between the control and TCDD-treated groups (17.1% versus 7.5%), there was no difference in the formation of AFL between the two groups in the *in vitro* metabolism assay. TCDD-treatment also did not affect the ability of the liver homogenates to bioactivate AFB₁ and AFM₁ *in vitro*. Unfortunately, enzyme activities could not be calculated in terms of P450 content, but rather only on a per mg protein basis, since zebrafish livers were too small to permit total P450 quantification. Analysis of CYP1A in the supernatant from zebrafish liver homogenates used in the metabolism and DNA adduction experiments verified that this protein and its associated EROD activity were induced by the TCDD treatment.

TCDD modulation of AFB₁-DNA adduction

The results from the investigation of TCDD modulation of hepatic AFB₁-DNA adduction showed an approximate 4-fold increase in the amount of hepatic DNA-adduction in the TCDD-treated group. A study by Walsh *et al.* (1992) also demonstrated that TCDD pretreatment increased DNA adduction and toxicity following exposure to AFB₁ in a human epidermal cell line. The increased binding in the TCDD-treated group of zebrafish cannot be explained by an increased ability to bioactivate AFB₁ to the reactive *exo*-8-9-epoxide, since the *in vitro* AFB₁ metabolism assay failed to show any difference in this activity between groups. This assay uses mouse cytosolic glutathione transferase which almost exclusively traps the *exo*-, but not the *endo*-AFB₁-

epoxide. The increased binding could be related to the increased production of AFM₁ and/or AFL-M₁, and further activation to a DNA binding species. The *in vitro* AFM₁ metabolism assay did demonstrate that zebrafish are quite capable of bioactivating AFM₁ to a reactive intermediate capable of binding glutathione and forming an adduct. *In vivo* genotoxicity tests of AFM₁ in *Drosophilia melanogaster* showed a only a 3-fold lower potency for damaging DNA compared to AFB₁ in one test, and equal genotoxicity in another test (Shibahara *et al.*, 1995). Work with rainbow trout hepatocytes also demonstrated that AFM₁ and AFL-M₁ had DNA binding values approximately 80% that of AFB₁ (Loveland, *et al.*, 1988).

In conclusion, zebrafish possess a CYP1A1-like protein that is readily inducible following i.p. administration or dietary exposure to β NF, but is refractory to Aroclor 1254 at the doses investigated in this study. Dietary exposure to TCDD, the most potent Ah receptor agonist, was also found to effective. Following exposure to TCDD, the metabolism of AFB₁ was altered in a pattern consistent with increases in CYP1A1, which is usually considered a detoxifying pathway. However, this pathway appears to correlate with an increase in the toxicity of AFB₁ in this species, as measured by an approximate 4-fold increase in hepatic AFB₁-DNA adduction.

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CHAPTER 4

CONCLUSIONS AND FUTURE DIRECTIONS

Zebrafish are efficient at metabolizing AFB, to both phase I and phase II metabolites, including bioactivation of AFB₁ to a reactive intermediate capable of binding to DNA, as shown by an in vitro AFB, metabolism assay and confirmed in the in vivo hepatic DNA-binding experiments. These data suggest that zebrafish should be fairly sensitive to the carcinogenic effects of AFB₁. However, Tsai¹ has demonstrated that zebrafish are actually resistant to AFB₁-induced carcinogenesis when the carcinogen is administered in the diet. Hepatic DNA-binding studies conducted following a dietary exposure to [3H]AFB, also showed only low levels of adducts (unpublished results). These results suggest a difference in absorbance and distribution of dietary AFB, compared to i.p. administration. It seems that if the carcinogen can reach the target organ, zebrafish are quite capable of bioactivating AFB₁. For future studies, this hypothesis could be evaluated by comparing the biodistribution of [3H]AFB, following gavage or injection, in order to measure the proportion of doses effectively delivered to the target organ by these two exposure routes. A logical companion study would be to compare the tumor response by dietary and i.p. administration of AFB₁ in zebrafish.

Zebrafish CYP1A responded to the common CYP1A inducers β NF and TCDD, as measured by catalytic activity and antibody detection of protein. CYP1A induction

¹Tsai, H.-W. (1996). Evaluation of zebrafish (*Brachydanio rerio*) as a model for carcinogenesis. [Ph.D. dissertation]. Oregon State University, Corvallis, OR.

occurred by 48 hours following i.p. administration of β NF. Surprisingly, there was no measurable response following dietary or i.p. administration of Aroclor 1254, a commercial mixture of PCBs that is an effective inducer in trout and other species. It would be interesting to further assess CYP1A response by measuring mRNA levels to see if transcription of this gene was activated or not. One could also explore the affects of other typical CYP1A inducers, such as 3-methylcholanthrene and some of the planar non-ortho substituted chlorinated biphenyls. By conducting investigations using more specific inducers, one could better evaluate if the agonist binding site might be more selective in this species.

The modulation of AFB₁ metabolism by a CYP1A inducer, in this case TCDD, was consistent with experiments conducted by other scientists. Due to increased levels of CYP1A, the metabolic profile shifted to formation of AFM₁ and AFL-M₁-glucuronide, the metabolite and its glucuronide conjugate typically associated with CYP1A. The 4-fold increase in hepatic DNA binding in the TCDD-pretreated group could not be explained by an enhanced capacity for bioactivation of AFB₁. It is postulated that AFM₁ is also bioactivated to a reactive intermediate that can bind to DNA. One approach to investigating this hypothesis is to essentially repeat the previous *in vivo* TCDD-modulation experiment, but this time going a step further. Instead of just measuring the total amount of radioactivity bound to DNA, one could look at the actual adduct profiles produced by the control or TCDD-pretreated groups following AFB₁ exposure. The adduct profile generated from the HPLC analysis should indicate whether AFM₁, or perhaps even AFL-M₁, forms an adduct with DNA, or if some other difference exists between the treatment groups in their DNA-adduction patterns.

The future of the zebrafish as a model for chemical carcinogenesis remains uncertain. Zebrafish do not appear to be sensitive to carcinogenesis following dietary exposures to AFB₁, N-nitrosodiethylamine and N-nitrosodimethylamine, but they did respond to the direct acting carcinogen, methylazoxymethanol acetate². As already mentioned, the lack of response in zebrafish following dietary exposures could be due to lack of absorption and distribution of the carcinogens. It should be noted that zebrafish do not have a stomach, and the effect of the absence of this organ on the process of absorption of compounds is not known. Other factors may play a significant role as well. For example, the dietary exposures could not be initiated until the fish were large enough to eat the diet used in carcinogen exposures. This timing typically coincided with sexual development², which in rainbow trout has correlated with decreasing sensitivity to carcinogens. Also, zebrafish are selective eaters, and it is difficult to assure that the proper dose has been given. In contrast to the lack of response of zebrafish following most dietary exposures of carcinogens, static water bath exposures of zebrafish fry and embryos resulted in tumorigenesis. Unfortunately, the metabolism of carcinogens other than AFB, has yet to be studied in zebrafish. Additional studies investigating the metabolic capabilities of this species could further elucidate the mechanistic basis for these contrasting results.

Another drawback of this fish model is that it is difficult to conduct biochemical studies in zebrafish. The liver is diffuse and difficult to remove, and large numbers of livers need to be pooled together in order to have enough tissue with which to work.

²Tsai, H.-W. (1996). Evaluation of zebrafish (*Brachydanio rerio*) as a model for carcinogenesis. [Ph.D. dissertation]. Oregon State University, Corvallis, OR.

In addition, due to the virtual impossibility of removing the liver intact, it appears that proteases are released during sampling, and if one is not careful, one can quickly lose biological activity of proteins. For a more detailed biochemical analysis, one might have to resort to using *in vitro* test systems, such as zebrafish cells in culture or cloned gene products. One will then have the additional question of the validity of the results as they relate to the *in vivo* system.

Because of the many difficulties and unexpected results encountered in the overall project to develop the zebrafish as a model for chemical carcinogenesis, many questions remain regarding zebrafish and their response to carcinogens.

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