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Chapter 3

Mycotoxins-Induced Oxidative Stress and Disease

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1. Introduction

Mycotoxins are pharmacologically active mold metabolites produced in a strain-specific way that elicit some complicated toxicological activities [1]. More than 300 secondary metabolites have been identified while only around 30 have true toxic properties [2]. The chemical structures of mycotoxins vary significantly, but they are low molecular mass organic compounds [3]. Mycotoxins are small and quite stable molecules which are extremely difficult to remove and enter the food and feed chain while keeping their toxic properties [4]. So, the occurrence of mycotoxins is regulated by legal limits in all developed countries [5]. Mycotoxin contamination of the feed and food is a global problem because more than 25% of world grain production is contaminated by mycotoxins [6]. The synthesis of mycotoxins by moulds is genetically determined and closely related to primary metabolic pathways, such as amino acid and fatty acid metabolism. However, the actual toxin production is modulated by environmental factors such as substrate composition and quality, humidity and temperature. The occurrence of mycotoxins in animal feed exhibits a geographic pattern, for example Aspergillus species meet optimal conditions only in tropical and subtropical regions, whereas Fusarium and Penicillium species are adapted to the moderate climate. Worldwide trade with food and feed commodities results in a wide distribution of contaminated material [7].

Plant selections for mycotoxin resistance have not created any significant results in protection against grain mycotoxins. The major problem comes from the fact that there are no safe levels of mycotoxins, because of synergistic interactions of many mycotoxins [2]. There is sufficient evidence from animal models and human epidemiological data to conclude that mycotoxins cause an important hazard to human and animal health [1]. The toxic effect of mycotoxins on animal and human health depends on the type of mycotoxin; level and duration of the exposure; age, health, and sex of the exposed individual, genetics,
dietary status, and interactions with other toxic insults. Thus, the severity of mycotoxin toxicity can be complicated by factors such as vitamin deficiency, caloric deprivation, alcohol abuse, and infectious disease [1,3,8].

Mycotoxins according to their chemical structure exert a broad variety of biological effects. The nature and intensity of these effects depend on the actual concentration of an individual mycotoxin and the time of exposure [7]. Cell proliferation of all mycotoxin treated blood mononuclear cells was significantly decreased at the highest concentrations of mycotoxins, but this decrease was significantly stronger for different mixtures of mycotoxins [9]. In addition, feed commodities are often contaminated with more than one mycotoxin, as mould species produce different mycotoxins at the same time. These co-occurring mycotoxins can exert additive effects, as for example various trichothecenes, but may also act antagonistically, as for example, observed with feeds containing trichothecenes and zearalenone, and commodities, containing aflatoxins and cyclopiazonic acid [7].

Mycotoxicoses are more common in underdeveloped countries and often remain unrecognized by medical professionals, except when huge numbers of people are involved [3]. In general, mycotoxin exposure is more likely to occur in parts of the world where poor methods of food handling and storage are common, where malnutrition is a problem, and where few regulations exist to protect exposed populations. The incidence of liver cancer varies widely from country to country, but it is one of the most common cancers in China, Philippines, Thailand, and many African countries. Worldwide, liver cancer incidence rates are 2 to 10 times higher in developing countries than in developed countries [10]. The occurrence of fumonisin B1 was correlated with the occurrence of a higher incidence of esophageal cancer in regions of Transkei (South Africa), China, and Northeast Italy [3]. In Africa and Asia where the occurrence of mycotoxins is common and a high percentage of the population is infected with hepatitis B or C mycotoxin reduction is obligatory [8]. One of the strategies for reducing the exposure to mycotoxins is to decrease their bioavailability by including various mycotoxins-adsorbing agents in the compound feed, which lead to reduction of mycotoxins uptake as well as distribution to the blood and target organs. Another strategy is the degradation of mycotoxins into non-toxic metabolites by using biotransforming agents such as bacteria/fungi or enzymes [4].

Diagnosis of animal mycotoxicosis is based on experimental studies with specific toxins and specific animals, very often under well-defined toxicological laboratory conditions, so that the results of such studies can be far from real-life or natural situations. Furthermore, factors such as breeding, sex, environment, nutritional status, as well as other toxic entities can affect the symptoms of intoxication and may contribute to the significance of mycotoxin damage on economic output and animal health [11]. The economic costs of mycotoxins are impossible to determine accurately [12], but the US Food and Drug Administration (FDA) estimated that in the US the mean economic annual cost of crop losses from the mycotoxins aflatoxins, fumonisins, and deoxynivalenol are $932 million USD [13]. While mycotoxin associated losses in industrial countries are typically market losses as a result of rejected crops, developing countries suffer additionally from health impacts [14]. Diagnosis is very
much dependent on receiving a sample of feed that was ingested prior to intoxication, but also on data from another representative group of animals of the facility and the results of a post-mortem examination [11, 13].

In the following table, the mycotoxins of major concern as feed contaminants are aflatoxins, ochratoxin A, Fusarium toxins (trichothecenes like, deoxynivalenol, diacetoxyscirpenol, nivalenol, T2-toxin/HT2-toxin, zearalenone and fumonisins) [4]. Moreover, the most predominant mycotoxigenic species in wheat grain were *A. flavus* with the ability to produce mycotoxins (aflatoxins B1, B2, G1 and G2 and sterigmatocystin) [15].

<table>
<thead>
<tr>
<th>Fungal species</th>
<th>Mycotoxin</th>
</tr>
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<tbody>
<tr>
<td><em>Aspergillus flavus</em>; <em>A. parasiticus</em></td>
<td>Aflatoxins</td>
</tr>
<tr>
<td><em>A. flavus</em></td>
<td>Cyclopiazonic acid</td>
</tr>
<tr>
<td><em>A. ochraceus</em>; <em>Penicillium viridicatun</em>; <em>P. cyclopium</em>; <em>P. expansum</em></td>
<td>Ochatoxin A</td>
</tr>
<tr>
<td><em>Fusarium culmorum</em>; <em>F. graminearum</em>; <em>F. sporotrichioides</em></td>
<td>Deoxynivalenol</td>
</tr>
<tr>
<td><em>F. sporotrichioides</em>; <em>F. poae</em></td>
<td>T-2 toxin</td>
</tr>
<tr>
<td><em>F. sporotrichioides</em>; <em>F. graminearum</em>; <em>F. poae</em></td>
<td>Diacetoxyscirpenol</td>
</tr>
<tr>
<td><em>F. culmorum</em>; <em>F. graminearum</em>; <em>F. sporotrichioides</em></td>
<td>Zearalenone</td>
</tr>
<tr>
<td><em>F. moniliforme</em></td>
<td>Fumonisins</td>
</tr>
<tr>
<td><em>Acremonium coenophialum</em></td>
<td>Ergopeptine alkaloids</td>
</tr>
</tbody>
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Table 1. The major toxigenic species of fungi and their mycotoxins [16]

**2. Route of mycotoxins exposure**

The most common route of exposure to mycotoxins is ingestion, but it may also involve dermal, respiratory, and parenteral routes, the last being associated with drug abuse [17]. In general, animals are directly exposed to mycotoxins through the consumption of mouldy feedstuffs, eating contaminated foods, skin contact with mould infected substrates and inhalation of spore-borne toxins [1]. Human exposure can be via one of two routes; direct exposure due to the consumption of mouldy plant products, or indirect exposure through the consumption of contaminated animal products containing residual amounts of the mycotoxin ingested by the food producing animals [18]. Human exposure to mycotoxins is further determined by environmental or biological monitoring. In environmental monitoring, mycotoxins are measured in food, air, or other samples; in biological monitoring, the presence of residues, adducts, and metabolites is assayed directly in tissues, fluids, and excretory products [19]. The risk of systemic toxicity resulting from dermal exposure increases in the presence of high toxin concentrations, occlusion, and vehicles which enhance penetration [20]. The main human and veterinary health load of mycotoxin exposure is related to chronic exposure [2].
3. Mycotoxins metabolism and induction of oxidative stress

Biodegradation of mycotoxins with microorganisms or enzymes is considered as the best strategy for detoxification of feedstuffs. This approach is considered as environmentally friendly approach in contrast to physicochemical techniques of detoxification. Ruminants are potential source of microbes or enzymes for mycotoxins biodegradation [21]. In vertebrate, mycotoxin is metabolized by cytochrome P450 enzymes to metabolite-guanine-N7 adduct (Fig 1). The carcinogenic potency is highly correlated with the extent of total DNA adducts formed in vivo [22].

Cytotoxicity and ROS generation are mechanisms of mycotoxins mediated toxicity. ROS are chemically reactive molecules containing oxygen. They are highly reactive due to the presence of unpaired electrons. ROS formed as a natural byproduct of the normal metabolism of oxygen have important roles in cell signaling and homeostasis. However, during times of environmental stress, ROS levels can increase dramatically as a result of oxidative stress [24]. Oxidative stress occurs when the concentration of ROS generated exceeds the antioxidant capability of the cell. In other words, oxidative stress describes various deleterious processes resulting from an imbalance between the excessive formation of ROS and limited antioxidant defenses [25]. Under normal conditions, ROS are cleared from the cell by the action of superoxide dismutase (SOD), catalase (CAT), or glutathione peroxidase (GPx). The main damage to cells results from the ROS-induced alteration of macromolecules such as polyunsaturated fatty acids in membrane lipids, proteins, and DNA. Additionally, oxidative stress and ROS can originate from xenobiotic bioactivation by prostaglandin H synthase (PHS) and lipoxygenases (LPOs) or microsomal P450s which can...
oxidize xenobiotics to free radical intermediates that react directly or indirectly with oxygen to produce ROS and oxidative stress [26] as in Fig (2). Moreover, the cell can tolerate a small to moderate amount of oxidative stress by producing antioxidant molecules e.g vitamin A, C &E and GSH and activates enzymes e.g. CAT, SOD, GPx, glutathione reductase (GR) and glutathione S transferase (GST) to counteract the excess oxidants [27]. LPO may bring about protein damage and inactivation of membrane-bound enzyme either through direct attack by free radicals or through chemical modification by its end products [28]. Reduction of cellular viability by mycotoxins was correlated with increases of ROS generation and MDA formation in concentration and time dependent manner [29]. The importance of oxidative stress and LPO in mycotoxins toxicity was confirmed by the protective effects of natural antioxidants [2]. Sporidesmin, the mycotoxin responsible for ‘facial eczema’ in ruminants, contains a disulphide group which appears to be intimately involved in its toxic action. The dithiol form of sporidesmin has been shown readily to undergo autoxidation in vitro in a reaction which generates superoxide radical (O_{2}^{−}) [30]. GST found in the cytosol and microsomes catalyzes the conjugation of activated aflatoxins with GSH, leading to the excretion of aflatoxin [31]. Variations in the level of the GST as well as variations in the cytochrome P450 system are thought to contribute to the differences observed in interspecies aflatoxin susceptibility [22, 32].

Figure 2. General pathways of ROS production and clearance [26]
4. Mycotoxin toxicity

The amount of mycotoxins needed to produce adverse health effects varies widely among toxins, as well as for each animal or person’s immune system. Two concepts are needed to understand the negative effects of mycotoxins on human health: *Acute toxicity*, the rapid onset of an adverse effect from a single exposure. *Chronic toxicity*, the slow or delayed onset of an adverse effect, usually from multiple, long-term exposures. Mycotoxins can be acutely or chronically toxic, or both, depending on the kind of toxin and the dose. Membrane-active properties of various mycotoxins determine their toxicity. Incorporation of mycotoxins into membrane structures lead to alterations in membrane functions. In general, mycotoxins effects on DNA, RNA, protein synthesis and the pro-apoptotic action (Fig. 3) causing changes in physiological functions including growth, development and reproduction [2]. Clinicians often arrange mycotoxins by the organ they affect. Thus, mycotoxins can be classified as hepatotoxins, nephrotoxins, neurotoxins, immunotoxins, and so forth. Cell biologists put them into generic groups such as teratogens, mutagens, carcinogens, and allergens [1].

![Diagram of mycotoxins affecting major sites in RNA and protein synthesis](image-url)

*Figure 3. Mycotoxins affecting major sites in RNA and protein synthesis [33]*
Aflatoxins

Aflatoxins occur in nuts, cereals and rice under conditions of high humidity and temperature. The two major *Aspergillus* species that produce aflatoxins are *A. flavus*, which produces only B aflatoxins, and *A. parasiticus*, which produces both B and G aflatoxins. Aflatoxins M1 and M2 are oxidative metabolic products of aflatoxins B1 and B2 produced by animals following ingestion, and so appear in milk, urine and faeces. Aflatoxicol is a reductive metabolite of aflatoxin B1. Aflatoxins are acutely toxic, immunosuppressive, mutagenic, teratogenic and carcinogenic compounds (classified as group 1 carcinogens according to the International Agency for Research on Cancer (IARC) [34]. Aflatoxins have been detected in the blood of pregnant women, in neonatal umbilical cord blood, and in breast milk in African countries, with significant seasonal variations [35]. The geographical and seasonal occurrence of aflatoxins in food and of kwashiorkor shows a remarkable similarity [36]. It has been hypothesized that kwashiorkor, a severe malnutrition disease, may be a form of pediatric aflatoxicosis [37]. Aflatoxins exposure accounts for about 40% of the load of disease in developing countries where a short lifespan is prevalent. Food systems and economics in developed country make the advance in aflatoxins management impossible [38]. The prevention of mycotoxins toxicity involves reduction of mycotoxin levels in foodstuffs and increasing the intake of diet components such as vitamins, antioxidants and substances known to prevent carcinogenesis [39]. The prevention of mycotoxin contamination of human foods could have a significant effect on public health in low-income countries due to enhanced food safety [40]. Chemoprotection against aflatoxins has been confirmed with the use of a number of compounds that either increase an animal’s detoxification processes [41] or prevent the production of the epoxide that leads to cytotoxicity [42]. For the animal feed industries, a major focus has been on developing food additives that provide protection from the mycotoxins. One approach has been the use of esterified glucomanoses and other yeast extracts that provide chemoprotection by increasing the detoxification of aflatoxin [41].

Figure 4. Metabolism of aflatoxin in liver [46]
After the absorption, highest concentration of the toxin is found in the liver [43]. Once in liver, aflatoxin B1 is metabolized by microsomal enzymes cytochrome P-450 3A4 to different metabolites through hydroxylation, hydration, demethylation and epoxidation. Variations in its catalytic activity of P-450 3A4 are important in issues of bioavailability and drug-drug interactions [44]. As in Fig (4) the hydroxylation of AFB1 at C4 or C22 produces, AFM1 and AFQ1, respectively. Hydration of the C2 – C3 double bond results in the formation of AFB2a which is rapidly formed in certain avian species [45]. AFP1 results from o-demethylation while the AFB1 – epoxide is formed by epoxidation at the 2,3 double bond. Aflatoxicol is the only metabolite of AFB1 produced by a soluble cytoplasmic reductase enzyme system [46].

The putative AFB1 epoxide is generally accepted as the active electrophilic form of AFB1 that may attack nucleophilic nitrogen, oxygen and sulphur heteroatoms in cellular constituents [47]. This highly reactive substance may combine with DNA bases such as guanine to produce alterations in DNA [36]. However, both humans and animals possess enzymes system, which are capable of reducing the damage to DNA and other cellular constituents caused by the 8,9-epoxide. For example GST mediates the conjugation reaction of the 8,9-epoxide to the endogenous compound GSH, this essentially neutralizes its toxic potential (Fig. 5). Animal species such as the mouse that are resistant to aflatoxin carcinogenesis have 3-5 times more GST activity than susceptible species such as the rat. Humans have less GST activity or 8,9-epoxide conjugation than rats or mice suggesting that humans are less capable of detoxifying this important metabolite [48].

![Figure 5. Biomarkers of aflatoxin exposure in an internal dose and a biologically effective dose. Biomarkers of exposure include aflatoxin M1, the internal dose includes the aflatoxin-mercapturic acid and aflatoxinalbumin adduct, and the biologically effective dose is reflected by the excretion of the aflatoxin-N7-guanine adduct formed by depurination leading to an apurinic (AP) site in DNA [49].](image-url)
The diseases caused by aflatoxin consumption are called aflatoxicosis. Acute aflatoxicosis results in death, however, chronic aflatoxicosis results in immune suppression and cancer [19]. Suppression of the cell-mediated immune response was mediated by altered cytokine expression [50]. Aflatoxins caused hepatotoxicity, nephrotoxicity and genotoxicity in somatic and germ cells, resulted in mitotic and meiotic delay in mice [51]. An increase in AFB1-8, 9-epoxide cause a significant increases in hepatic LPO level [52]. Peroxidation of membrane lipids initiated loss of membrane integrity; membrane bound enzyme activity and cell lysis [53]. LPO was significantly increased in the liver, kidney [54] and testis [55] of aflatoxin-treated mice as compared to controls. However, GSH levels declined significantly in the liver, kidney and testis after 45 days of aflatoxin treatment [56]. Moreover, AFB1 intake and expression of enzymes involved in AFB1 activation/detoxification may play an important role in hepatitis B virus-related hepatocarcinogenesis [57]. The results of a clinical trial suggest that chlorophyllin may have a role in preventing dietary exposure to aflatoxin B1 by reducing its oral bioavailability [58].

Strategies for reducing exposure and risk from aflatoxin in developing countries should be carefully tested and validated using clinical trial designs with biomarkers serving as objective endpoints. Clinical trials and other interventions are designed to translate findings from human and experimental investigations to public health prevention. Both primary (to reduce exposure) and secondary (to alter metabolism and deposition) interventions can use specific biomarkers as endpoints of efficacy. In a primary prevention trial, the goal is to reduce exposure to aflatoxins in the diet. A range of interventions includes planting pest-resistant varieties of staple crops, attempting to lower mould growth in harvested crops, improving storage methods following harvest, and using trapping agents that block the uptake of unavoidably ingested aflatoxins. In secondary prevention trials, one goal is to modulate the metabolism of ingested aflatoxin to enhance detoxification processes, thereby reducing internal dose and subsequent risk [49] (Fig. 6).

Ochratoxin A (OTA)

Ochratoxins are the first major group of mycotoxins identified after the discovery of the aflatoxins [59]. OTA is found in a variety of plant food products such as cereals. Because of its long half life, it accumulates in the food chain [60]. OTA is absorbed passively throughout the gastrointestinal tract and actively in the kidneys. Highest amounts of OTA could be found in the blood and it is distributed in kidney, liver, muscle and adipose tissue in a decreasing order. The toxin is excreted primarily in the urine, and to a lesser degree in bile and also in milk. The half-life of experimentally orally ingested OTA is shorter than intravenously injected OTA [61]. According to IRAC [34] OTA is classified as group 1 carcinogens. Structure-activity studies suggested that the toxicity of OTA may be attributed to its isocoumarin moiety and lactone carbonyl group. OTA reduces the expression of several genes regulated by nuclear factor-erythroid 2 p45-related factor (Nrf2) and reduces the expression of antioxidant enzymes through inhibition of Nrf2 [62, 63]. OTA toxicity may be involved in the development of certain kidney diseases through generation of oxidative stress [64]. Chronic administration of low dose of OTA caused morphological and functional
changes in proximal tubules and administration of date extract protects against OTA-induced tubule’s tissue damage [65]. However, antioxidant treatment failed to prevent the development of OTA-induced tumors in animal models [66]. Indomethacin and aspirin were found to prevent OTA genotoxicity in the urinary bladder and kidney of mice [67]. OTA causes acute depletion of striatal dopamine and its metabolites, accompanying evidence of neural cell apoptosis in the substantia nigra, stratum and hippocampus [68].

Figure 6. Strategies for reducing exposure and risk from aflatoxin in developing countries [49]

OTA has complex mechanisms of action that include oxidative stress, bio-energetic compromise, mitochondrial impairment, inhibition of protein synthesis, production of DNA single strand breaks and formation of OTA-DNA adducts [69-71]. OTA induced renal toxicity and carcinogenicity may be mediated by an Nrf2-dependent signal transduction pathway [63]. It is a mitochondrial poison causing mitochondrial damage, oxidative burst, LPO and interferes with oxidative phosphorylation [72, 73]. OTA was found to chelate ferric ions (Fe³⁺), facilitating their reduction to ferrous ions (Fe²⁺), which in the presence of oxygen, provided the active species initiating LPO [74]. OTA hydroquinone/ quinone couple was generated from the oxidation of OTA by electrochemical, photochemical and chemical processes resulting in redox cycling and in the generation of ROS [75]. OTA impairs the antioxidant defense of the cells making them more susceptible to oxidative damage [62] and a reduction in cellular antioxidant defense may contribute to the production of OTA-dependent oxidative damage [76].

Studies carried out in several countries including Tunisia, Egypt and France, have indicated a link between dietary intake of OTA and the development of renal and urothelial tumours [77-81]. OTA is known to affect the immune system in a number of mammalian species. The type of immune suppression experienced appears to be dependant on a number of factors,
including the species involved, the route of administration, the doses tested, and the methods used to detect the effects [82]. OTA causes immunosuppression following prenatal, postnatal and adult-life exposures. These effects include reduced phagocytosis and lymphocyte markers [83] and increased susceptibility to bacterial infections and delayed response to immunization in piglets [9]. OTA induces apoptosis in a variety of cell types in vivo and in vitro that mediated through cellular processes involved in the degradation of DNA [84]. Moreover, the immunosuppressant activity of OTA is characterized by size reduction of vital immune organs, such as thymus, spleen, and lymph nodes, depression of antibody responses, alterations in the number and functions of immune cells, and modulation of cytokine production (TNF-α and II-6). The immunotoxic activity of OTA probably results from degenerative changes and cell death following necrosis and apoptosis, in combination with slow replacement of affected immune cells, due to inhibition of protein synthesis [85]. Finally, it is proposed that a network of interacting epigenetic mechanisms, including protein synthesis inhibition, oxidative stress and the activation of specific cell signalling pathways is responsible for OTA carcinogenicity [86] (Fig. 7).

Figure 7. Scheme to illustrate the oxidative stress-mediated mode of action proposed for OTA. Increased production of ROS, RNS, and RCS is likely to originate either from direct redox reactions involving OTA or through the inhibition of cellular defenses such as through the inhibition of transcription factors as Nrf2 which regulates enzymes with antioxidant properties. The generation of radicals will induce macromolecular oxidative damage such as oxidized DNA bases which may be converted into mutation resulting into generation of transformed cells [66].

**Trichothecenes**

Trichothecenes (TCs) are mycotoxins produced mostly by members of the *Fusarium* genus and other genera (e.g. *Trichoderma, Trichothecium, Myrothecium* and *Stachybotrys*). By now,
more than 180 different trichothecenes and trichothecene derivatives have been isolated and characterized [87, 88]. TCs were found in air samples collected during the drying and milling process on farms, in the ventilation systems of private houses and office buildings and on the walls of houses with high humidity [89-90]. They can be divided into four categories according to both their chemical properties and their producer fungi;

1. **Type A**: functional group other than a ketone at C8 position (e.g.; T-2, HT-2, DAS);
2. **Type B**: carbonyl functions at C8 position (e.g.; DON, NIV, FUS-X, 3-acetyl-deoxynivalenol, 15-acetyl-deoxynivalenol);
3. **Type C**: second epoxide group at C7, 8 or C9, 10; (e.g.; crotocin and baccharin);
4. **Type D**: macrocyclic ring system between C4 and C15 with two ester linkages (e.g.; satratoxin G, H, roridin A and verrucarin A) [87, 91, 92].

TCs exposure leads to apoptosis both *in vitro* and *in vivo* in several organs such as lymphoid organs, hematopoietic tissues, liver, intestinal crypts, bone marrow and thymus [91, 93]. Acute high dose toxicity of TCs is characterized by diarrhea, vomiting, leukocytosis, haemorrhage, and circulatory shock and death, whereas chronic low dose toxicity is characterized by anorexia, reduced weight gain, neuroendocrine and immunologic changes [91, 94]. The myelotoxicity was considered highest for T-2 and HT-2 toxins and lowest for DON and NIV [94]. TCs are toxic to many animal species, but the sensitivity varies considerably between species and also between the different TCs [93]. Cellular effects on DNA and membrane integrity have been considered as secondary effects of the inhibited protein synthesis. The toxin binds to the peptidyl transferase, which is an integral part of the 60S ribosomal subunit of mammalian ribosome. TCs interfere with the metabolism of membrane phospholipids and increase liver LPO *in vivo*. Also, some TCs are shown to change the serotonin activity in the central nervous system, which is known to be related in the regulation of food intake [88].

**T-2 Toxin**

T-2 toxin is a cytotoxic secondary fungal metabolite that belongs to TCs family produced by various species of *Fusarium* (*F. sporotrichoides*, *F. poae*, *F. equiseti*, and *F. acuminatum*), which can infect corn, wheat, barley and rice crops in the field or during storage [95]. T-2 toxin is a well known inhibitor of protein synthesis through its high binding affinity to peptidyl transferase resulting in trigger of ribotoxic stress response that activate mitogen-activated protein kinases [96]. Moreover, T-2 toxin interferes with the metabolism of membrane phospholipids and increases liver LPO [97]. Also, T-2 toxin suppresses drug metabolizing enzymes such as GST [98]. T-2 toxin treated mice showed a time-dependent increase in ROS generation, depletion of GSH, increases in LPO and protein carbonyl content in the brain. Moreover, the gene expression profile of antioxidant enzymes showed a significant increase in SOD and CAT via the dermal route and GR and GPx via the subcutaneous route [99]. General signs of T-2 include nausea, emesis, dizziness, chills, abdominal pain, diarrhea, dermal necrosis, abortion, irreversible damage to the bone marrow, reduction in white blood cells and inhibition of protein synthesis [100, 101]. Moreover, the effects of T-2 toxins on the immune system include changes in leukocyte counts, delayed hypersensitivity,
depletion of selective blood cell progenitors, depressed antibody formation and allograft rejection [39]. Also, T-2 toxin has a direct lytic effect on erythrocytes [102]. T-2 toxin can induce apoptosis in many types of cells bearing rapid rates of proliferation [103] and increased the expression of both oxidative stress and apoptosis related genes in hepatocytes of mice [104]. T-2 toxin induces neuronal cell apoptosis in the fetal and adult brain [68]. In this aspect, it suggested that dysfunction of the mitochondria and oxidative stress might be the main factor behind the T-2 toxin-induced apoptosis in the fetal brain [105]. ROS activate caspase-2 which play a crucial role in the control of apoptosis [106, 107]. Moreover, it was demonstrated that T-2 toxin induced cytotoxicity in HeLa cells is mediated by generation of ROS leading to DNA damage and trans activation of p53 protein expression which leads to shift in the ratio of Bax/Bcl-2 in favour of apoptosis and subsequent release of Cyt-c from mitochondria followed by caspase cascade [99].

**Fumonisins**

Fumonisins produced by the fungus *Fusarium verticillioides*, a widespread fungal contaminants of various cereals, predominantly corn [2, 108]. Fumonisin B1 (FB1) and B2 are of toxicological significance, while the others (B3, B4, A1 and A2) occur in very low concentrations and are less toxic [3]. FB1 is poorly absorbed and rapidly eliminated in feces. Minor amounts are retained in liver and kidneys. FB1 does not cross the placenta and is not teratogenic in vivo in rats, mice, or rabbits, but is embryotoxic at high, maternally toxic doses [109]. FB1 has been linked to a number of diseases in humans and animals [1, 40]. FB1 increases oxidative DNA damage, as measured by increased DNA strand breaks and malondialdehyde adducts in rat liver and kidney in vivo [111]. As shown in Fig. (9) an alternative mechanism of action of FB1 involves the disruption of the de novo sphingolipid biosynthesis pathway by inhibition of the enzyme ceramide synthase [68, 112]. The inhibition of sphingolipid biosynthesis disrupts numerous cell functions and signaling pathways, including apoptosis and mitosis, thus potentially contributing to carcinogenesis through an altered balance of cell death and replication [113]. Disruption of sphingolipid metabolism leads to changes in the sphinganine to sphingosine ratio [114] as demonstrated in rat liver and mouse kidney at carcinogenic doses of FB1 [115].

FB1-induced DNA damage and hepatocarcinogenesis in experimental models can be modulated by a variety of factors including nutrients, chemopreventive agents, and other factors such as food restriction and viral infection, as well as genetic polymorphisms [118]. In rat C6 glioma cells, FB1 inhibits protein synthesis, causes DNA fragmentation and cell death, increases 8-hydroxy-2’-deoxyguanosine, induces LPO, and cell cycle arrest [119]. Also, the signs of apoptosis were increased caspase-3 like protease activity and internucleosomal DNA fragmentation [120]. Furthermore, the disruption of membrane structure, the enhancement of membrane endocytosis, and the increase in membrane permeability caused by FB1 in macrophages provide additional insight into potential mechanisms by which the fumonisins might enhance oxidative stress and cellular damage [121].
Figure 8. Pathway of de novo synthesis (not in boxes) and turnover of sphingolipids (boxed) in animal cells, and their inhibition by fumonisins. Fumonisins inhibit the synthesis of ceramides by specifically binding to sphingosine and sphinganine [116, 117]

Zearalenone

Zearalenone (ZEA) is produced mainly by *Fusarium graminearum* and related species, principally in wheat and maize. ZEA and its derivatives produce estrogenic effects in various animal species [3]. The structure of the ZEA similar to steroids and binds to ER as an agonist. There are two major biotransformation pathways for ZEA in animals (1) hydroxylation catalyzed by 3α- and 3β- hydroxysteroid dehydrogenase (HSDs) resulting in the formation of α- and β-ZOL; (2) conjugation of ZEA and its metabolites with glucuronic acid, catalyzed by uridine diphosphate glucuronyl transferase. Consequently, ZEA is a substrate for 3α-HSD and 3β-HSD present in many steroidogenic tissues, such as liver, kidney, testis, prostate, hypothalamus, pituitary, ovary, intestine [122, 123]. The adverse effects of ZEA are partly determined by the processes of elimination, because the biliary excretion and entero-hepatic cycling are important processes affecting the fate of ZEA and explaining a different sensitivity between animals [124]. α and β Zearalenol metabolites caused cytotoxicity by inhibiting cell viability, protein and DNA syntheses and inducing oxidative damage and over-expression of stress proteins. However, the ZEA metabolites exhibited lower toxicity than ZEA, with β zearalenol being the more active of the two metabolites [125]. In addition, oxidative damage is likely to be evoked as one of the main pathways of ZEA toxicity which may initiate event at least in part contribute to the mechanism of ZEA induced genotoxic and cytotoxic effects [126]. ZEA and its derivatives compete effectively with 17 β-E2 for the specific binding sites of the oestrogen receptors (ERs) occurring in different organs. Two subtypes of ER exist, ER-α and ERβ that are differently distributed in the body. Binding of ZEA and its derivatives initiate a sequence of
events known to follow estrogen stimulation of target organs [127, 128]. So, the effect of ZEA and its metabolites depends upon the reproductive status (prepubertal, cycling or pregnant) of the affect animals [123]. ZEA do not induce apoptosis in porcine ovaries [129], however, apoptosis is the principal mechanism contributing to germ cell depletion and testicular atrophy following ZEA exposure [130]. Moreover, ZEA has potent effects on the expression of chicken splenic lymphocytes cytokines at the mRNA level [131].

**Patulin**

Patulin (PAT) is a toxic chemical contaminant produced by several species of mould, especially within *Aspergillus, Penicillium* and *Byssochlamys*. It is the most common mycotoxin found in apples and apple-derived products such as juice, cider, compotes and other food intended for young children. Exposure to this mycotoxin is associated with immunological, neurological and gastrointestinal outcomes [132]. PAT-induced nephropathy and gastrointestinal tract malfunction have been demonstrated in animal models [133]. The toxic effects of PAT on various cells related to its activity on SH groups [134]. Moreover, it suggested that PAT-induced apoptosis is mediated through the mitochondrial pathway without the involvement of p53 [12]. The interaction with sulfhydryl groups of macromolecules by PAT and the subsequent generation of ROS plays a primary role in the apoptotic process. The genotoxic effects might be related to its ability to react with sulfhydryl groups and to induce oxidative damage [135]. PAT was found to reduce the cytokine secretion of IFN-γ and IL-4 by human macrophages [136] and of IL-4, IL-13, IFN-γ, and IL-10 by human peripheral blood mononuclear cells and human T cells [137]. The clinical signs of PAT toxicosis are typical of the nervous syndrome. Animals present hyperaesthesia, lack of coordination of motor organs, problems with ingestion and digestion. At the molecular level, PAT alters ion permeability and/or intracellular communication, causing oxidative stress and cell death (116).

**Citrinin**

Citrinin is a toxic metabolite produced by several filamentous fungi of the genera *Penicillium, Aspergillus* and *Monascus*, which has been encountered as a natural contaminant in grains, foods, feedstuffs, as well as biological fluids. Some analytical systems have been developed for its detection and quantification [138]. As one of mycotoxins, citrinin possesses antibiotic, bacteriostatic, antifungal and antiprotozoal properties. While it is also known as a hepato-nephrotoxin in a wide range of species [139], in vitro studies have demonstrated that citrinin produced multiple effects on renal mitochondrial function and macromolecule biosynthesis that ultimately resulted in cell death [140]. Citrinin inhibited the oxygen consumption rate by about 45 % and inhibited the glucose utilization of BHK-21 cells by about 86 % due to alterations in mitochondrial function and in the glycolytic anaerobic pathway [141]. Citrinin occurred frequently together with another nephrotoxin–ochratoxin A in foodstuffs such as cereals, fruits, meat [142] and cheese [143]. Citrinin can act synergistically with ochratoxin A to depress RNA synthesis in murine kidneys [144, 145]. The simultaneous exposure of rabbits to citrinin with OTA even at sub-clinical dietary levels potentiated the OTA induced nephrotoxicity at ultrastructural level [146]. To avoid the
direct/indirect intake of citrinin, it is important to develop detoxification methods for citrinin during food processing. So far, there have been several reports on the detoxification of citrinin. The investigation on thermal decomposition and detoxification showed that, in the presence of a small amount of water, heating citrinin at 130°C caused a significant decrease in its toxicity to Hela cells [147]; whereas heating at 150°C in water caused formation of highly toxic compounds [148].

**Ergot Alkaloids**

The ergot alkaloids are among the most fascinating of fungal metabolites. They are classified as indole alkaloids and are derived from a tetracyclic ergoline ring system [149]. These compounds are produced as a toxic cocktail of alkaloids in the sclerotia of species of Claviceps, which are common pathogens of various grass species. Ergotism is still an important veterinarian problem. The principal animals at risk are cattle, sheep, pigs, and chickens. Clinical symptoms of ergotism in animals include gangrene, abortion, convulsions, suppression of lactation and hypersensitivity [150]. More recently, pure ergotamine has been used for the treatment of migraine headaches. Other ergot derivatives are used as prolactin inhibitors, in the treatment of Parkinsonism, and in cases of cerebrovascular insufficiency [149]. The therapeutic administration of ergot alkaloids may cause sporadic cases of human ergotism [151]. Ergotism is extremely rare today, primarily because the normal grain cleaning and milling processes remove most of the ergot so that only very low levels of alkaloids remain in the resultant flours. In addition, the alkaloids that are the causative agents of ergotism are relatively labile and are usually destroyed during baking and cooking [3].

**Satratoxin G**

Satratoxin G is one of the most potent macrocyclic TCs produced by Stachybotrys chartarum [152]. Roridin A is a commercially available macrocyclic TC used as a stratoxin G substitute, and roridin L2 is a putative biosynthetic precursor of satratoxin [153]. Satratoxin G is potent inhibitors of eukaryotic translation that are potentially immunosuppressive. It rapidly binds small and large ribosomal subunits in a concentration- and time-dependent manner that was consistent with induction of apoptosis [154]. A signal transduction pathway in satratoxin-induced apoptosis in HL-60 cells involves, caspase-3 activation through activation of both caspase-8 and caspase-9 along with cytosolic release of cytochrome c and fragmentation of nucleosomal DNA by DFF-40/CAD [155].

**Roridin E**

Roridin E is a well-known macrocyclic trichothecene mycotoxin possessing potent anti-proliferative activity against cancer cell lines [156]. Four new isolated from a marine-derived fungus, Myrothecium roridum strain 98F42 [157]. One of them, 12-deoxy derivative of roridin E, showed reduced cytotoxicity about 80-fold less than that of roridin E against human promyelocytic (HL-60) and murine leukemia (L1210) cell lines [158]. Treatment of rats with roridin E caused minimal toxicity on the hepatic and renal tissue, however, coadministration of linoleic acid with roridin E resulted in increase toxicity due to increased incorporation to the cell membrane or inhibit its biotransformation [159].
5. Mycotoxins and apoptosis

Apoptosis is a process for maintenance of tissue homeostasis. Several processes, such as initiation of death signals at the plasma membrane, expression of pro-apoptotic oncoproteins, activation of death proteases and endonucleases combine to cause cell termination. ROS may play a major role in apoptosis. GSH depletion increases the % of apoptotic cells [160]. In general, apoptosis is considered as a common mechanism of toxicity of various mycotoxins [68]. TCs induce apoptosis response via mitochondrial and non-mitochondrial mechanisms [161]. The amphophilic nature of TCs facilitates their cytotoxic effect on cell membranes and inside the cell interact with ribosome and mitochondria causing inhibition of protein synthesis [162]. FB₁ and OTA are able to induce apoptosis and necrosis in porcine kidney PK15 cells [163]. This is because the structure of FB₁ resembles sphingoid bases which regulate cell growth, differentiation, transformation and apoptosis, and so it is not surprising that FB₁ can alter growth of certain mammalian cells. The involvement of the TNF signal transduction pathway in FB₁ induced apoptosis in African green monkey kidney fibroblasts has been shown [164]. Moreover, TNF-α production is responsible for FB₁ induced apoptosis in mice primary hepatocytes [165]. Over expression of cytochrome P450-sensitized hepatocyte to TNF-α-mediated cell death was associated with increased LPO and GSH depletion [166]. FB₁ was reported to increase induction of cytochrome P450 isoforms and caused peroxidation of membrane lipids in isolated rat liver nuclei as well as GSH depletion of in pig kidney cells [167-169]. GSH depletion is known to activate c-Jun N-terminal kinase through redox inhibition of GST, which normally binds to and inhibits stress kinases [170]. Stimulation of apoptosis and necrosis in porcine granulosa cells by ZEA is dose-dependent manner via a caspase-3- and caspase-9-dependent mitochondrial pathway [171]. At the molecular level, fumonisins inhibit ceramide synthase and disrupt sphingolipid metabolism therefore influence apoptosis and mitosis [109]. The immunotoxic activity of OTA probably results from degenerative changes and cell death following necrosis and apoptosis in combination with slow replacement of affected immune cells due to inhibition of protein synthesis [85]. Moreover, PAT induce DNA damage and cell cycle arrest along with intrinsic pathway mediated apoptosis which may have dermal toxicological implications [172].

Satratoxin H is thought to induce apoptosis of PC12 cells through the activation of p38 mitogen activated protein kinase and c-Jun N-terminal kinase in GSH-sensitive manner [173]. Chemoprotective effects of flavonoid compounds against aflatoxins were confirmed in hens [174]. Moreover, cysteine and GSH has protective effect against PAT in the incident of rumen microbial ecosystem, however vitamin C and ferulic acid did not demonstrate an effect [175]. Metallothioneins (MTs) are four major isoforms found in cytoplasm, lysosomes, mitochondria and nuclei of mammalian cells [176]. MT-1 and 2 have ubiquitous tissue distribution particularly in liver, pancreas, intestine, and kidney, whereas MT-3 is found in brain and MT-4 in skin [177]. MT can play important role in the process of mycotoxins detoxification probably by redistribution of significant ions to transcriptional factors and interactions with oxygen radicals that may be generated by mycotoxins [23]. Nivalenol, a
trichothecene mycotoxin induces apoptosis in HL60 cells and that intracellular calcium ion plays a role in the nivalenol-induced secretion of IL-8 from this cell line [178].

6. Mycotoxins as therapeutics compound

Cumulative knowledge about toxins structure and mechanism of action, as well as recent progress in the fields of cell biology, immunology, molecular biology and nanotechnology, enabled the development of different targeting strategies that are vital for converting a lethal toxin into a therapeutic agent. Fig. 9 showed three targeting strategies in toxin based therapy.

i- Ligand targeted toxins upon administration to patients are internalized and intoxicate diseased cells, sparing healthy cells that do not display the target on their surface. ii- protease activated toxins: the toxin is engineered to be cleaved and activated by a disease-related intracellular or extracellular protease. Toxin cleavage may enhance cell-binding and/or translocation, stabilization or catalytic activity of the toxic moiety specifically in protease expressing cells, leading to their suppression. iii- toxin based suicide gene therapy: a DNA construct, encoding for a toxic polypeptide whose expression is regulated by a specific transcription regulation element, is delivered to a heterogeneous cell population [179].

![Figure 9. Three targeting strategies in toxin based therapy [179].](image)

Because of their pharmacological activity, some mycotoxins or mycotoxin derivatives have found use as antibiotics, growth promoters, and other kinds of drugs. Ergocryptine is an ergot alkaloid that affects dopaminergic activity principally by interacting with D2-type receptors [180]. The bromation derivative has increased dopamine agonist activity, and is used against Parkinsonism and to reduce growth hormone secretion and milk production [181]. Ergotamine was among the most effective available agents for relieving migraine attacks [182]. Ergometrine and the semi-synthetic methylergometrine have been widely used for the prevention and treatment of excessive uterine bleeding following birth and also to initiate delivery [183]. Lysergic acid diethylamide is a serotonin receptor agonist [184] and can also interact with dopamine receptors to make it a useful tool for probing the
biochemical basis for behaviour [185]. Methysergide a semi-synthetic ergot alkaloid is a serotonin antagonist used in the treatment of migraine and is used for daily preventive therapy rather than in acute cases [184]. The TCs have been associated with various biological properties, such as antiviral especially as inhibitors of the replication of Herpes, antibiotic, antimalarial, antileukemic and immunotoxic [59,186]. Mycoestrogenic zearalenone is suspected to be a triggering factor for central precocious puberty development in girls. Due to its chemical resemblance to some anabolic agents used in animal breeding, ZEA may also represent a growth promoter in exposed patients [187]. Development of cyclosporine A as immunosuppressive drug has been traced back to the stimulus derived from the first highly-active cyclopeptides from *Amanita mushrooms* [188].

7. Conclusion

Mycotoxins are produced in a strain-specific way, and elicit some complicated and overlapping toxigenic activities in sensitive species that include carcinogenicity, inhibition of protein synthesis, immunosuppression, dermal irritation, and other metabolic perturbations. Mycotoxins usually enter the body via ingestion of contaminated foods, but inhalation of toxigenic spores and direct dermal contact are also important routes. There is sufficient evidence from animal models and human epidemiological data to conclude that mycotoxins pose an important danger to human and animal health. Trichothecenes cause protein synthesis inhibition via binding to the 18s rRNA of the ribosomal large subunit as a major mechanism underlying induction of cell apoptosis. T-2 toxin triggers a ribotoxic response through its high binding affinity to peptidyl transferase which is an integral part of the 60 s ribosomal subunit and interferes with the metabolism of membrane phospholipids and increases liver lipid peroxides. SH is thought to induce caspase-3 activation and apoptosis through the activation of MAPK and JNK in a GSH-sensitive manner. FB1-induced inhibition of ceramide synthesis can result in a wide spectrum of changes in lipid metabolism and associated lipid-dependent pathways. OTA has complex mechanisms of action that include mitochondrial impairment, formation of OTA-DNA adducts and induction of oxidative stress and apoptosis through caspase activation. Accordingly, the strict control of food quality, in both industrialized and developing countries, is therefore necessary to avoid mycotoxicosis.

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8. References


