REVIEW ARTICLE

Fumonisins: oxidative stress‑mediated toxicity and metabolism in vivo and in vitro

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Abstract Fumonisins (FBs) are widespread Fusarium toxins commonly found as corn contaminants. FBs could cause a variety of diseases in animals and humans, such as hepatotoxic, nephrotoxic, hepatocarcinogenic and cytotoxic effects in mammals. To date, almost no review has addressed the toxicity of FBs in relation to oxidative stress and their metabolism. The focus of this article is primarily intended to summarize the progress in research associated with oxidative stress as a plausible mechanism for FB-induced toxicity as well as the metabolism. The present review showed that studies have been carried out over the last three decades to elucidate the production of reactive oxygen species (ROS) and oxidative stress as a result of FBs treatment and have correlated them with various types of FBs toxicity,

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indicating that oxidative stress plays critical roles in the toxicity of FBs. The major metabolic pathways of FBs are hydrolysis, acylation and transamination. Ceramide synthase, carboxylesterase FumD and aminotransferase FumI could degrade $FB₁$ and $FB₂$. The cecal microbiota of pigs and alkaline processing such as nixtamalization can also transform $FB₁$ into metabolites. Most of the metabolites of $FB₁$ were less toxic than $FB₁$, except its partial (pHFB₁) metabolites. Further understanding of the role of oxidative stress in FB-induced toxicity will throw new light on the use of antioxidants, scavengers of ROS, as well as on the blind spots of metabolism and the metabolizing enzymes of FBs. The present review might contribute to reveal the toxicity of FBs and help to protect against their oxidative damage.

Keywords Fumonisins · Oxidative stress · DNA damage · Lipid peroxidation · Reactive oxygen species · Antioxidants · Metabolism · Metabolizing enzymes

U-118MG Human U-118MG glioblastoma ZEN Zearalenone

Introduction

Fumonisins (FBs), belong to the relatively recently discovered group of mycotoxins mainly produced by *Fusarium verticillioides* and *Fusarium proliferatum*, are widespread mycotoxins commonly found as corn contaminants (Marschik et al. [2013](#page-18-0)). *A. niger* is also a source of fumonisins, such as Fumonisin₂ (FB₂), Fumonisin₄ (FB₄) and Fumoni- \sin_6 (FB₆) (Mansson et al. [2010](#page-18-1)). In addition, FBs have also been found in wheat, barley, sorghum, rice, wine, raisins, black tea leaves, asparagus, pine nuts and Thai coffee beans (Logrieco et al. [2010;](#page-18-2) Marin et al. [2007](#page-18-3); Martins et al. [2001;](#page-18-4) Mateo and Jimenez [2000](#page-18-5); Nielsen et al. [2015](#page-19-0)). They are structurally characterized by an eicosane backbone, esterified with two tricarballylic acid groups (Fig. [1](#page-2-0)). Fumonisin₁ (FB₁) and FB₂ are the most important representatives of FBs (European Commission [2003](#page-17-0)). In most cases from field samples, FB_1 , FB_2 and Fumonisin₃ (FB_3) constituted approximately 70, 20 and 10 % of FBs, respectively (Howard et al. [2002\)](#page-18-6).

FBs are found all around the world. In Hebei Province of China during 2011–2013, the incidence of FBs $(FB_1 + FB_2)$ for all samples was 46.4 % (Li et al. [2015b](#page-18-7)). In Shandong Province of China in 2014, the incidences of FB_1 , FB_2 and FB_3 in the corn products were 98.1 %, and the simultaneous occurrence of FB_1 , FB_2 and FB_3 was observed in 76.7 % of the corn products (Li et al. [2015a](#page-18-8)). The incidence of FBs in nine provinces of China was reported to be from 81.0 to 10.3 % in maize samples in 2012 (Fu et al. 2015). A recent study revealed that FB₂ was also detected in 37.5 % of the onion samples from the Taif region in Saudi Arabia (Gherbawy et al. [2015\)](#page-17-2). In animal feeds distributed in South Korea in 2011, the contamination rate of feed ingredients with FB_1 and FB_2 was 50 and 40 %, respectively (Seo et al. [2013\)](#page-19-1). FB₂ was found in 29 % of the raisin samples at levels ranging from 7.1 to 25.5 µg/kg in raisins from Western Greece regions in 2011 (Perrone et al. [2013](#page-19-2)). In Iran, the natural occurrences of FB_1 , FB_2 and FB_3 were 68.2, 42.6 and 31.7 %, respectively (Sorensen et al. [2009](#page-19-3)). A prevalence survey in Taiwan showed that there was a contamination rate of 40 % in domestic animal feeds, and the average contaminated level was 4.5 mg/kg (Cheng et al. [2006](#page-17-3)).

FBs could cause a variety of diseases in animals and humans, such as hepatotoxic, nephrotoxic, hepatocarcinogenic and cytotoxic effects in mammals (Dall'Asta et al. [2009](#page-17-4)). FB₁ is the most abundant of the naturally occurring analogues, and it was classified as possibly carcinogenic to humans by the International Agency for Research

on Cancer (IARC [2002\)](#page-18-9). It was reported that purified $FB₁$ could cause both liver and kidney tumors in rodents in long-term feeding studies (Creppy [2002\)](#page-17-5). The high incidence of human esophageal cancer was reported to be associated with FBs, such as in Cixian Country of Hebei Province in China (Wang et al. [2000\)](#page-20-0), and in the former Transkei region of the Eastern Cape Province of South Africa (Rheeder et al. [1992\)](#page-19-4). $FB₁$ was reported to inhibit embryonic sphingolipid synthesis and has been implicated as a risk factor for neural tube defects (NTDs) in cultured mouse embryos (Sadler et al. [2002](#page-19-5)) and in the sensitive LM/Bc mouse (Gelineau-van Waes et al. [2005](#page-17-6), [2009;](#page-17-7) Marasas et al. [2004](#page-18-10); Sadler et al. [2002](#page-19-5); Voss et al. [2009](#page-19-6)). Furthermore, $FB₁$ could cause cardiotoxicity and pulmonary edema in swine associated with *F. verticillioides* exposure (Colvin and Harrison [1992](#page-17-8); Smith et al. [1999](#page-19-7)). Horses were the more susceptible species to FBs, and the main mycotoxicosis of horses is leukoencephalomalacia caused by the FB₁ and FB₂ produced by *Fusarium spp.* (Riet-Correa et al. [2013](#page-19-8); Voss et al. [2007](#page-19-9)).

FBs could inhibit ceramide synthases with the subsequent disruption of sphingolipid metabolism and sphingolipid-dependent physiological processes (Gelineau-van Waes et al. [2009;](#page-17-7) Voss et al. [2009](#page-19-6); Zitomer et al. [2009](#page-20-1)). $FB₁$ modulates immunity in animals, and it has immunocytotoxic effects by increasing IL-10 and IL-4 mRNA levels in the spleen of Balb/c mice together with the decrease in mRNA levels of IFN- γ and TNF- α compared with controls (Abbes et al. 2015). Similarly, FB₁ could significantly inhibit the expression of IL-1β, IL-2, IFN- α and IFN-γ when broilers were exposed to 15 mg FB₁ per kg for 3 weeks (Cheng et al. 2006). Together with AFB₁ or not, $FB₁$ increased levels of IL-4 and decreased levels of IL-10 in spleen mononuclear cells (SMC) in both in vivo and in vitro studies (Theumer et al. [2002,](#page-19-10) [2003\)](#page-19-11). Moreover, recent studies indicated that the generation of reactive oxygen species (ROS) and oxidative stress was closely related to the immunotoxic effect elicited by the mycotoxins (Kotan et al. [2011;](#page-18-11) Mary et al. [2012;](#page-18-12) Theumer et al. [2010](#page-19-12)). There has been considerable focus for the last several years on the potential of $FB₁$ to induce oxidative stress. In vitro studies conducted on several cell lines have suggested the possible role of oxidative stress on FB_1 -induced cytotoxicity and apoptosis (Bernabucci et al. [2011;](#page-17-10) Domijan et al. [2015](#page-17-11); Ferrante et al. [2002](#page-17-12); Kang and Alexander [1996](#page-18-13); Klaric et al. [2007](#page-18-14); Kouadio et al. [2005](#page-18-15); Mary et al. [2012](#page-18-12); Mobio et al. [2003;](#page-19-13) Stockmann-Juvala et al. [2004a](#page-19-14), [b](#page-19-15)). Similar results were also observed in mice (Abbes et al. [2015](#page-17-9)), rats (Hassan et al. [2014\)](#page-18-16) and broiler chicks (Poersch et al. [2014](#page-19-16)). Oxidative stress will occur in cells when the concentration of ROS exceeds the antioxidant capacity (Mishra et al. [2014\)](#page-18-17). High concentrations of ROS lead to the dis-Fig. 1 Chemical structures of the main fumonisins **Fig.** 1 Chemical structures of the main fumonisins lipid peroxidation in the lipid membrane causing damage to the cell membrane, and DNA damage by propagating a chain reaction (Mishra et al. [2014](#page-18-17); Zuo et al. [2015](#page-20-2)). In the toxicity of FBs, oxidative stress and ROS might play important roles in the induction of FB-induced damage to lipids, DNA and proteins. To date, several reviews have been published on the determination method of FBs, biomarkers of FB-induced toxicity, apoptosis and carcinogenesis (Arranz et al. [2004;](#page-17-13) Scott [2012](#page-19-17); Shephard et al. [2007](#page-19-18); Stockmann-Juvala and Savolainen [2008](#page-19-19)). Although oxidative stress was briefly mentioned in one of the reviews 7 years ago (Stockmann-Juvala and Savolainen [2008](#page-19-19)), some new articles about the role of oxidative stress induced by FBs have been published. Therefore, it is necessary to review the new progress of the toxic effects on oxidative stress induced by FBs. The scope of this review is primarily intended to summarize the evidence associated with a role of oxidative stress as a plausible mechanism for FBinduced toxicity. The studies related to FBs toxicity under in vitro and in vivo conditions are summarized in Tables [1](#page-4-0) and [2](#page-6-0), respectively. Furthermore, the metabolizing routes, metabolizing enzymes, factors affecting the metabolism of FBs and the toxicity of metabolites of FBs were also reviewed. The present review shows that over the past two decades, oxidative stress and ROS generation might correlate closely with various types of toxicity, as well as with the metabolism of FBs, and the antioxidant mechanism contributed greatly to finding more effective antioxidants to inhibit FBs-induced toxicity.

Oxidative stress and toxicity

Generation of reactive oxygen, oxidative stress and reactive nitrogen species

Oxidative stress is initiated by ROS such as perhydroxy radical (HOO⁻), hydroxyl radical (HO[']) and superoxide anion (O_2^-) , and by reactive nitrogen species (RNS) including nitric oxide (Mishra et al. [2014\)](#page-18-17).

Mary et al. ([2012\)](#page-18-12) documented that when SMC from Wistar rats were exposed to 10 μ M FB₁ for up to 48 h, the treatment increased total ROS. The mitochondrial complex I, CYP450, the arachidonic acid metabolism and the NADPH oxidase system were involved in the ROS generation induced by the FB_1 group (Mary et al. [2012](#page-18-12)). In cell cultures of rat primary astrocytes and human neuroblastoma (SH-SY5Y), FB_1 (0.5, 5 and 50 μ M), as a neurodegenerative mycotoxin, decreased the rate of mitochondrial and cellular respiration, and increased ROS production in mitochondria by inhibiting mitochondrial complex I (Domijan and Abramov [2011](#page-17-14)). However, the effect of $FB₁$ on ROS over-production is a consequence rather than a mechanism of its toxicity (Domijan and Abramov [2011](#page-17-14); Galvano et al. 2002). FB₁ increased the production of ROS in human U-118MG glioblastoma (U-118MG) cells, showing significant dose- and time-level effects after incubation with 100 μ M FB₁ for 48–144 h and with 10 μ M FB₁ for 72 and 144 h (Stockmann-Juvala et al. [2004b](#page-19-15)). After cells were treated with FB₁ (0.1–100 μ M) for 0–144 h, FB₁ caused a dose-dependent increase in ROS production in mouse C6 glioblastoma and rat GT1-7 hypothalamic cells, but was without any effect in human SH-SY5Y cells (Stockmann-Juvala et al. [2004a\)](#page-19-14). It was interesting to note that ROS production was observed in SH-SY5Y cells by Domijan and Abramov, whereas it was not noted by Stockmann-Juvala (Domijan and Abramov [2011;](#page-17-14) Stockmann-Juvala et al. [2004a](#page-19-14)). However, to date, there has been no generation of RNS reported in the study of $FB₁$. When birds were fed diets containing FB_1 (5, 10 and 15 mg/kg) for 3 weeks, $15 \text{ mg } FB_1/kg$ feed had no effect on inducible nitric oxide synthase (iNOS) (Cheng et al. [2006\)](#page-17-3).

FBs‑mediated oxidative damage

Oxidative stress up-regulates the antioxidant defense system and resulted in damage to cellular macromolecules, such as DNA, lipids and proteins. Subsequently, oxidative stress led to cell death by apoptotic or necrotic mechanisms. During the progress, DNA damage, enhanced lipid peroxidation and protein damage occurred (Tables [1](#page-4-0), [2](#page-6-0)). The schematic representation of FBs induced the damage to DNA, lipids and proteins is shown in Fig. [2](#page-8-0).

Damage to lipids

Oxidative damage to lipids of the cell membrane increases lipid peroxidation. Lipid peroxidation can be measured by monitoring the changes in thiobarbituric acid-reacting substances (TBARS) and malondialdehyde (MDA). As one part of TBARS, MDA was the most abundant individual aldehyde resulting from lipid peroxidation, and its level is a marker of lipid oxidation (Klaric et al. [2007;](#page-18-14) Pirinccioglu et al. [2010](#page-19-20); Zhou et al. [2015\)](#page-20-3).

FBs could significantly increase lipid peroxidation. $FB₁$ (35 and 70 µg/mL) significantly increased MDA production in bovine peripheral blood mononuclear cells (Bernabucci et al. [2011](#page-17-10)). The exposure of SMC from Wistar rats to $FB₁$ (10 µM) for 48 h significantly raised the formation of MDA relative to the control (Mary et al. [2012\)](#page-18-12). Applied for 48 h, only the highest dose of FB_1 (5 µg/mL) significantly increased TBARS (57 %) (Klaric et al. [2007\)](#page-18-14). Compared with other Fusarium toxins such as deoxynivalenol (DON) and zearalenone (ZEN), $FB₁$ showed the highest production of MDA in Caco-2 cells, identifying that $FB₁$ causes the highest level of oxidative damage to lipids (Kouadio et al.

Table 1 In vitro fumonisin-related oxidative stress studies

protection against FB₁ toxicity

protection against FB₁ toxicity

Table 1 continued

 2005). MDA production induced by $FB₁$ was also noted in the study of oxidative stress in SH-SY5Y neuroblastoma, rat C6 glioblastoma, mouse GT1-7 hypothalamic cells, C6 glioma cells (with normal p53 status) and p53-null mouse embryonic fibroblasts, indicating that lipid peroxidation induced by FB_1 might be a common phenomenon (Mobio et al. [2003;](#page-19-13) Stockmann-Juvala et al. [2004a](#page-19-14)). A similar report showed that the treatment of FB_1 (5 and 10 μ M) for 24 h increased MDA production in the macrophage cell line J774A.1 (Ferrante et al. [2002](#page-17-12)). An in vivo study also found that FB_1 (100 µg/kg b.w.) resulted in a significant increase in MDA level in liver and kidney tissues (Hassan et al. 2014). After feeding a diet contaminated with $FB₁$ (100 mg/kg) for 21 days, hepatic TBARS levels in male broiler chicks were also increased (Poersch et al. [2014](#page-19-16)). Theumer et al. [\(2010](#page-19-12)) reported that after exposure to experimental diets contaminated with a culture material containing 100 mg/kg of $FB₁$ for 90 days, the MDA levels were increased in SMC (Theumer et al. [2010](#page-19-12)). The level of MDA in serum was elevated in SD rats treated with $FB₁$ (100 µg/ kg b.w.) by gavage for 30 days (Sun et al. [2006](#page-19-21)). Abel and Gelderblom [\(1998](#page-17-16)) reported that dietary $FB₁$ levels of 250 and 500 mg $FB₁/kg$ increased the level of TBARS significantly in the liver of rats fed FB_1 over 21 days. Also, in the 250 mg FB_1/kg group, subcellular fractionation of the liver of the rats showed a significant increase in TBARS in the microsomes. Furthermore, the level of TBARS was increased in a dose-dependent manner, associated with an increase in cytotoxicity in primary rat hepatocytes (Abel and Gelderblom [1998](#page-17-16)). Another study revealed that female Sprague–Dawley (SD) rats fed FB-contaminated diet (200 mg/kg diet) showed a significant increase in MDA compared with the control group (Hassan et al. [2010](#page-18-19)). However, MDA concentration did not change from control after the treatment with FB_1 (5, 50 and 500 µg/kg b.w.) for 4, 24 and 48 h in the liver of male Wistar rats. The low $FB₁$ dose and short exposure time were suggested as the possible reasons (Domijan et al. [2008](#page-17-17)). In contrast, MDA level in the liver of male SD rats was significantly increased when the rats were fed FB-contaminated diet (200 mg/kg diet) for 3 weeks, indicating that a sufficient exposure time of FBs could also lead to lipid peroxidation, even at a low concentration of FBs (El-Nekeety et al. [2007](#page-17-18)). In a study to determine whether very low doses of $FB₁$ induce oxidative stress in the rat kidney and liver, rats were treated orally with FB_1 (200 ng/kg b.w. and 50 µg/kg b.w.). The results found that FB_1 did not affect MDA concentration in the liver, while both $FB₁$ doses increased MDA concentration in the kidney, suggesting that the kidney was more sensitive to the toxicity of $FB₁$ (Domijan et al. [2007](#page-17-19)).

extract, *PGE* panax ginseng extract, *LAB* lactic acid bacteria

Fig. 2 Oxidative stress-mediated mode of action proposed for fumonisins. The increased generation of ROS as well as a decrease in antioxidant status may induce lipid, protein and DNA oxidation, leading to toxicity and apoptosis via ERK and the intrinsic mitochondrial pathway. The heat shock proteins might be induced by oxidative stress and protect cells against oxidative damage

Damage to DNA and RNA

The threat of oxidative damage is particularly significant to DNA. DNA damage can be measured by alkaline comet (SCGE) assay, cytokinesis-block micronucleus (CBMN) assay and Fpg-modified comet assay (Domijan et al. [2006,](#page-17-20) [2015](#page-17-11)). The formation of the major oxidative DNA damage product 8-hydroxydeoxyguanosine (8-OH-dG) was also used as an indicator of oxidative DNA damage (Ihsan et al. [2011](#page-18-22)).

Domijan revealed that FB_1 (20 μ g/mL) already induced DNA damage in the primary human peripheral blood lymphocytes (HPBLs) after 1-h exposure (Domijan et al. [2015](#page-17-11)). An 8-h exposure of SMC to the FB₁ (10 μ M) significantly increased the 8-OH-dG levels, compared with the control (Mary et al. [2012](#page-18-12)). In a study to compare the cytotoxic effect of DON, ZEN and FB_1 (1–150 μ M) and the pathways leading to cell death and related to oxidative stress in Caco-2 cells, DNA synthesis is inhibited by $FB₁$ with the highest IC50 of 20 μ M among three kinds of toxins (Kouadio et al. [2005](#page-18-15)). DNA fragmentation was studied in U-118MG cells exposed to 10–100 μ M FB₁ for 144 h, and $FB₁$ increased inter-nucleosomal DNA fragmentation as the formation of DNA ladders (Stockmann-Juvala et al. [2004b](#page-19-15)). Mobio et al. [\(2003](#page-19-13)) reported that the treatment with $FB₁$ (3 and 36 μ M) for 24 h induced a significant and doserelated increase in 8-OH-dG and DNA fragmentation in both C6 glioma and MEF cells. The maximum DNA breakage was already observed at 9 μ M of FB₁ in both C6 glioma and MEF cells. However, MEF cells showed a higher sensitivity to $FB₁$ even at lower $FB₁$ concentrations. In addition, FB_1 (9 and 18 μ M) incubation significantly increased the rate of deoxycytosine methylation in C6 glioma cells but not in MEF cells, suggesting that different cells showed different sensitivity to FBs (Mobio et al. [2003](#page-19-13)). DNA fragmentation in the liver and kidney was increased after female SD rats were exposed to FB_1 (100 µg/kg b.w.) for 12 weeks (Hassan et al. [2014](#page-18-16)). Another study showed that the DNA damage was found in 13.7 and 81.7 % (comet assay) and in 2.8 and 7.0 % (micronucleus technique) in control and 100 mg/kg $FB₁$ groups in a subchronic study in male Wistar rats, respectively (Theumer et al. [2010](#page-19-12)). Similarly, another study suggested that FB_1 -induced micronucleated polychromatic erythrocytes (Mn-PCEs) in bone marrow increased DNA and RNA in the liver after female SD rats were fed an FB-contaminated diet (200 mg/kg diet) for 4 weeks (Hassan et al. [2010\)](#page-18-19).

The role of oxidative stress in FB_1 -induced DNA damage has been illustrated by Domijan et al. [\(2008](#page-17-17)) in male adult Wistar rats treated with single $FB₁$ doses (5, 50 and 500 µg/kg b.w.) and killed at 4, 24 and 48 h after treatment. There was a significant increase in the DNA mobility of liver cells after 48 h following treatment with 50 and 500 µg/kg b.w. compared to the control. Moreover, tail intensity appeared to be a more sensitive parameter for detecting DNA damage, even at 5 µg/kg b.w. after 48 h. This study proved that FB_1 -induced DNA damage is timeand dose-dependent, and Wistar rats were sensitive to $FB₁$ (Domijan et al. [2008\)](#page-17-17). To investigate DNA damage induced by FB₁, rats orally received FB₁ (0.2, 50 and 500 μ g/ kg b.w., respectively) for 5 days. The results showed that even at the lowest dose of $FB₁$, the tail length, tail intensity and olive tail moment (OTM) obtained with the standard comet assay and Fpg-modified comet assay were significantly higher in treated animals than in controls. Furthermore, the Fpg-modified comet assay showed significantly greater tail length, tail intensity and OTM in all treated animals than in the standard comet assay, indicating that oxidative stress is likely to be responsible for DNA damage induced by $FB₁$. However, DNA damage detected by the standard comet assay at all $FB₁$ doses indicated that some other mechanism was also involved (Domijan et al. [2006](#page-17-20)). Additionally, treatment with $FB₁$ led to splenic and hepatic DNA fragmentation in 85 % of the test animals after fasted

rats were injected intravenously with a single dose of $FB₁$ at 1.55 mg/kg b.w. into the tail vein, suggesting that oxidative damage caused by FB_1 may be one of the underlining mechanisms of FB_1 -induced cell injury and DNA damage (Atroshi et al. [1999\)](#page-17-21).

Damage to proteins

Apart from lipids and DNA, the possible major target of oxidative damage is proteins that can be transformed into protein carbonyls (PCs) (Domijan et al. [2007\)](#page-17-19). Mary et al. identified that the 48-h incubation of SMC with $FB₁$ (10 µM) significantly raised the carbonyl levels compared to the control (Mary et al. [2012\)](#page-18-12). The inhibition of protein synthesis has been reported in a few studies. In one study to compare the inhibition of protein synthesis among DON, ZEN and $FB₁$ in Caco-2 cells, IC50 of 5, 8.8 and 19 μ M for DON, FB₁ and ZEN identified that protein synthesis is not a specific target of $FB₁$. The toxic effects of $FB₁$ act through lipid peroxidation rather than protein synthesis (Kouadio et al. 2005). In the $FB₁$ concentration range (3–30 µM), protein synthesis was inhibited in both C6 glioma (IC50 = 6 μ M) and MEF cells (IC50 = 5 μ M) after 24-h incubation with a superimposable kinetic (Mobio et al. [2003\)](#page-19-13). Domijan et al. [\(2007](#page-17-19)) reported that $FB₁$ (200 ng/kg b.w. and 50 µg/kg b.w.) could cause oxidative damage of proteins in the kidney of male Wistar rats. $FB₁$ did not affect PCs concentrations in the liver, while in the kidney, both $FB₁$ doses increased the concentration of PCs, indicating that proteins are also more susceptible to oxidative lesions in the kidney than in the liver (Domijan et al. [2007](#page-17-19)).

Alterations in antioxidant status

The enzymatic antioxidant defense systems are responsible for protecting from ROS such as O_2^- , HO^{\cdot} and hydrogen peroxide $(H₂O₂)$ (Yang and Lee [2015](#page-20-4)). The primary antioxidant enzymes are SOD, CAT, glutathione reductase (GR) and GPx. These antioxidant enzymes can serve as redox biomarkers because they are the first to indicate the antioxidant state through oxidation/reduction processes (Yang and Lee [2015](#page-20-4)). GSH is the most abundant intracellular antioxidant involved in the protection of cells against oxidative damage and in various detoxification mechanisms (Shi et al. [2015](#page-19-23)).

The antioxidant enzymes were regarded as important mediators of the immunotoxicity induced by $FB₁$. After Balb/c mice were treated daily for 2 weeks by oral gavage with FB_1 (100 µg/kg), the results showed that in the spleen, exposure to mycotoxins led to an increase in lipid peroxidation, IL-10 and IL-4 mRNA levels, but decreased GSH content and down-regulated expression of GPx and SOD,

and of IFN- γ and TNF- α mRNA (Abbes et al. [2015](#page-17-9)). FB₁ might significantly increase intracellular reactive oxygen metabolites (ROM) and decrease both cytoplasmic SOD and GPx mRNA abundance, suggesting that FB_1 might induce cytotoxicity through impairment of the oxidative status of bovine peripheral blood mononuclear cells (PBMCs) (Bernabucci et al. [2011\)](#page-17-10).

The activities of antioxidant enzymes in oxidative stress are implicated as a mechanism of $FB₁$ neurotoxicity (Domijan and Abramov [2011](#page-17-14); Stockmann-Juvala et al. [2004a](#page-19-14)). After 24-h treatment of FB_1 (0.5–200 μ M), the level of GSH in astrocytes was significantly increased in SH-SY5Y cells (Domijan and Abramov [2011](#page-17-14)). The ability of $FB₁$ to increase GSH level was explained by its direct action on the proteins responsible for GSH synthesis or/and by its mild ROS production that can stimulate GSH synthesis (Dinkova-Kostova and Talalay [2008](#page-17-22); Domijan and Abramov [2011](#page-17-14); Galvano et al. [2002\)](#page-17-15). However, long-term exposure of U-118MG cells to $FB₁$ resulted in decreased GSH levels. Significant depletion of GSH was noted after incubation with 1, 10 and 100 μ M FB₁ for 144 h and after incubation with 100 μ M FB₁ for 48 and 72 h (Stockmann-Juvala et al. $2004b$). Similarly, $FB₁$ decreased GSH levels in mouse C6 glioblastoma and rat GT1-7 hypothalamic cells but had no effects in human SH-SY5Y cells (Stockmann-Juvala et al. [2004a](#page-19-14)). It was interesting to note that the trend of GSH level induced by $FB₁$ was increased in SH-SY5Y cells, as reported by Domijan and Abramov, whereas it was not affected in the study of Stockmann-Juvala (Domijan and Abramov [2011;](#page-17-14) Stockmann-Juvala et al. [2004a](#page-19-14)). The difference in the trend of GSH level in both studies might be related to the different concentrations and incubation time of $FB₁$.

The misbalance of the enzymatic antioxidant defense systems was noted in vivo when animals were administered with FB_1 . Abbes et al. ([2015\)](#page-17-9) found that FB_1 decreased GSH content and down-regulated the expression of GPx and SOD in spleen when Balb/c mice were treated daily for 2 weeks by oral gavage with FB_1 (100 μ g/kg) (Abbes et al. [2015](#page-17-9)). Similarly, female SD rats receiving $FB₁$ (100 μ g/kg b.w.) for 12 weeks decreased GSH content in the liver and kidney and down-regulated the gene expression of antioxidants enzymes such as GPX, SOD and CAT (Hassan et al. [2014](#page-18-16)). When broiler chicks were administered the $FB₁$ diet (100 mg/kg) for 21 days, the hepatic CAT activity in the liver was significantly increased, showing that subacute exposure of $FB₁$ induced liver oxidative stress in broiler chicks (Poersch et al. [2014](#page-19-16)). In SMC of male Wistar rats subchronically (90 days) fed with culture material containing 100 mg/kg of $FB₁$, the CAT and SOD activities were increased in the poisoned animals, indicating the hypothesis of an oxidative stress-mediated genotoxicity induced by FB_1 (Theumer et al. [2010\)](#page-19-12). FB_1 showed hepatotoxicity

in C57BL/6 N mice accompanied by decreased GSH in the liver following the subcutaneous injection of 2.25 mg $FB₁/$ kg b.w. once daily for 5 days (He et al. [2006\)](#page-18-21). Intravenous injection of the tail vein with FB_1 (1.55 mg/kg) markedly decreased GSH levels in the liver (20 % of the control) and spleen (26 % of the control) (Atroshi et al. [1999\)](#page-17-21). However, in the research of early toxic effects of $FB₁$ in rat liver, GSH concentration did not change when male adult Wistar rats were treated with single $FB₁$ doses (5, 50 and 500 µg/ kg b.w.) and killed at 4, 24 and 48 h after treatment. The reason for this was suggested to be the low $FB₁$ dose and short exposure time (Domijan et al. [2008\)](#page-17-17). However, with a longer exposure time of FB_1 study, FB_1 did not affect the activity of CAT and SOD in the liver and kidney when rats were treated orally with FB_1 (200 ng/kg b.w. and 50 µg/kg b.w.) for 5 days (Domijan et al. [2007\)](#page-17-19). Rumora et al. ([2007\)](#page-19-22) noted that cellular GSH content in the liver of rats exposed for 2 days did not differ from control animals when Wistar rats were treated intraperitoneally (i.p.) with $FB₁$ (0.5 mg/ kg b.w.) for 2 or 7 days, whereas in the liver of rats treated with $FB₁$ for 7 days, GSH concentrations increased significantly (24 % increase), and kidney GSH levels in animals treated for 2 and 7 days decreased significantly (15 and 27 % decrease, respectively); this indicates that the enzymatic antioxidant defense systems in the kidney seemed to be more sensitive to oxidative stress than that in the liver (Rumora et al. [2007](#page-19-22)).

Stress‑mediated biological response

Oxidative stress plays important roles in a large number of biological responses and cell signaling pathways. Thus, the significant changes in gene expression and the stimulation or inhibition of signal transduction usually result in many toxicological effects. The role of FB-mediated oxidative stress in the induction of apoptosis and their respective cell signaling pathways has been studied widely in different cell systems (Tables [1,](#page-4-0) [2\)](#page-6-0).

Apoptosis and cell signaling

Apoptosis is regarded as a common result of oxidative stress caused by ROS production, disturbed GSH defense system and lipid peroxidation (Higuchi [2003\)](#page-18-23). The caspasedependent pathway was usually stimulated and resulted in cell apoptosis (Stockmann-Juvala et al. [2004b\)](#page-19-15). It was suggested that human U-118MG glioblastoma cells treated with $FB₁$ underwent caspase-dependent apoptosis. The highest dose of FB_1 (100 µM) caused a significant increase in caspase-3-like protease activity in cells at 12, 48 and 72 h of exposure; caspase-3-like protease activity was not affected at other time points (24 and 144 h) or with lower concentrations of FB_1 (1 and 10 μ M), indicating that the

activation of caspase-3 might be one of the events causing an increase in ROS production, and subsequent lipid peroxidation and reduction in intracellular GSH levels (Stockmann-Juvala et al. [2004b](#page-19-15)). Abbes et al. [\(2015](#page-17-9)) reported that increased caspase-3 activity was noted in the spleens of rats after exposure to FB_1 (100 µg/kg) daily for 2 weeks by oral gavage (Abbes et al. [2015\)](#page-17-9).

The p53 gene is known to suppress cell cycle progression in response to DNA damage, thus preventing the transmission of damaged genetic information from one cell generation to the next (Yin et al. [1997](#page-20-5)). An interesting study found that the suppression of cell cycle progression was observed in C6 glioma but not in p53-null MEF cells incubated with $FB₁$, indicating that cells lacking mechanisms governed by the p53 gene would be more susceptible to neoplastic cascade or mutation following DNA lesions induced by this mycotoxin (Mobio et al. [2003](#page-19-13)).

The perturbations of the cellular redox state or the generation of ROS stimulate redox-sensitive signaling molecules, such as mitogen-activated protein kinases (MAPKs) and heat shock proteins (Hsps). The MAPK superfamily of serine/threonine kinases is activated by a number of extracellular stimuli and is involved in signal transduction cascades that play an important regulatory role in cell apoptosis (Matsuzawa and Ichijo [2005](#page-18-24)). Three major mammalian MAPKs were extracellular signal-regulated kinase (ERK), c-Jun N-terminal kinase (JNK) and p38 MAPK (Wang et al. [2012](#page-20-6); Wu et al. [2014\)](#page-20-7). The expression of Hsp25/27 and Hsp70 might affect cell survival by interfering with apoptotic machinery and inhibiting apoptosis both upstream and downstream of the caspase cascade in a caspase-independent manner. These proteins could also decrease ROS production, affect cellular GSH levels and neutralize the toxic effects of oxidized proteins (Escobedo et al. [2004](#page-17-23); Ferns et al. 2006). FB₁ could also increase MAPKs activation and the expression of Hsp25 and Hsp70 in the liver and kidney of male Wistar rats given 0.5 mg FB₁/kg b.w. intraperitoneally for 2 or 7 days. Phospho-JNK signal was activated even when cells were treated with $FB₁$ for 2-day treatment. However, $FB₁$ increased Hsp25 expression and decreased Hsp70 expression in the rat kidney after a 7-day treatment with $FB₁$. These findings documented that $FB₁$ could lead to tissue-specific activation and expression of redox-sensitive signaling molecules, and kidney cells were more sensitive to adverse effects of $FB₁$ (Rumora et al. [2007](#page-19-22)).

 $Ca²⁺$ homeostasis of the cells is one of the many important functions. The proliferation of cells, to undergo apoptosis, induction of oxidative stress and physiological functions such as signal transduction, is part of Ca^{2+} homeostasis (Cig and Naziroglu [2015](#page-17-25)). A recent study revealed that FB₁ (0.5, 5 and 50 μ M) resulted in mitochondrial injury and decreased calcium uptake in SH-SY5Y cells and led to the sustained deregulation of calcium homeostasis and

presumably cell death (Domijan and Abramov [2011](#page-17-14)). $FB₁$ (1.55 mg/kg b.w.) administration caused total Ca^{2+} in liver to increase within 4 h (204 % of control), while pretreatment with CoQ10 (30 mg CoQ10/kg diet) together with ^l-carnitine (2.8 mg carnitine/kg diet), alpha-tocopherol (30 IU vitamin E/kg diet) and selenium (1 mg selenium as sodium selenite/kg diet) could decrease the activities of Ca^{2+} in the liver (Atroshi et al. [1999\)](#page-17-21).

Prevention of FB‑mediated oxidative stress

Various compounds, including free radical scavengers, have been shown to combat FB-induced cell damage (Fig. [3\)](#page-12-0). As an antioxidant, sodium copper chlorophyllin (CHL) (0.1–100 µg/mL) exhibited free radical scavenging activity in the DPPH assay and protected against cell and DNA damage induced by FB_1 (20 μ g/mL) (Domijan et al. [2015\)](#page-17-11). Co-treatment using lactic acid bacteria (LAB) (2 \times 10⁹ CFU/L, approximately 2 mg/kg) with $FB₁$ (100 µg/kg) suppressed levels of DNA fragmentation, normalized splenic lipid peroxidation and increased GSH levels, up-regulated the expression of GPx and SOD and normalized mRNA levels of the analyzed cytokines (Abbes et al. [2015](#page-17-9)). Panax ginseng extract (PGE) exerted antioxidant and anti-apoptotic effects, which resulted from its ability to play a role in increasing the antioxidant status as well as lowering the oxidative damage of nucleic acids in the body (Abdel-Wahhab et al. [2010\)](#page-17-26). PGE contains many classes of compounds, including salicylic acid, caffeic acid, vitamins C and E, which have antioxidant activity and are important to enhance the non-enzymatic antioxidants of the body such as vitamins C and E, and the antioxidant capacity including the enzymatic antioxidants such as CAT, GPx, GSH and SOD (Ramesh et al. [2012a](#page-19-24), [b\)](#page-19-25). Research has shown that PGE (20 mg/kg b.w.) combined treatments with $FB₁$ (100 µg/kg b.w.) could suppress DNA fragmentation in the liver of female SD rats, normalize lipid peroxidation and increase GSH in the liver and kidney and up-regulate the expression of GPx, SOD1 and CAT mRNA (Hassan et al. [2014](#page-18-16)).

In another study, the ethanol extract of *Aquilegia vulgaris* L. (5 and 10 mg/kg b.w.) was identified to be against the oxidative stress and the genotoxicity in female SD rats fed with FB-contaminated diet (200 mg/kg diet). *A. vulgaris* L. (Ranunculaceae) is a perennial herb indigenous in central and southern Europe, Africa and Asia (Hassan et al. [2010\)](#page-18-19). The FBs within corn powder consisted of 68 % FB_1 , 21 % FB₂ and 11 % FB₃ based on the total FBs in the corn powder. The combined treatment showed significant improvements in GSH, total antioxidant capacity and MDA and succeeded to normalize PCEs, NCEs, PCEs/NCEs ratio, DNA and RNA in the liver and histological pictures in the liver tissues (Hassan et al. [2010](#page-18-19)). It was interesting to

note that different teas showed varying effects on the hepatotoxicity, cancer promotion and activities of CAT, GPx and GR, as well as GSH status in the rat liver. Unfermented rooibos and honeybush significantly reduced the total number of foci, respectively, while fermentation seems to reduce the protective effect of the herbal teas (Marnewick et al. [2009](#page-18-20)). A study showed that royal jelly (RJ) has a protective effect against FBs toxicity, and this protection was dose-dependent. Co-treatment with FBs (200 mg/kg diet) plus RJ (100 or 150 mg/kg b.w.) resulted in a significant improvement in the histological and histochemical pictures of the liver and kidney, and all of the tested parameters such as ALT, AST, triglycerides, cholesterol, HDL, LDL, creatinine, MDA, GPx and SOD and uric acid levels (El-Nekeety et al. [2007\)](#page-17-18). Mobio et al. ([2003\)](#page-19-13) reported that vitamin E

Fig. 3 Schematic illustration of preventive effect of different compounds including antioxidants and free radical scavengers on fumonisins-induced oxidative stress. The use of different antioxidants such as lactic acid bacteria (LAB), panax ginseng extract (PGE) and the ethanol extract of *Aquilegia vulgaris* L. suppressed levels of DNA fragmentation, improved the total antioxidant status and led to the prevention of apoptosis. Sodium copper chlorophyllin (CHL) exhibited free radical scavenging activity and protected against DNA damage induced by $FB₁$. Similarly, royal jelly (RJ) has a protective effect against FB toxicity by decreasing lipid oxidation as well as improving antioxidant status. The mixture of CoQ10 together with L-carnitine, alpha-tocopherol and selenium decreased DNA damage and the activities of Ca²+, ASAT and ALAT in the liver, thus leading to the prevention of apoptosis

(25 mM) inhibited MDA production in both C6 glioma and p53-null MEF cells, while transresveratrol (3 mM) and epicatechin (15 mM) decreased MDA formation in C6 glioma cells (Mobio et al. [2003\)](#page-19-13). The prevention of DNA damage and cell injury has also been reported by Atroshi et al. (1999) (1999) in FB₁-treated rats $(1.55 \text{ mg/kg b.w.})$. The study demonstrated that pretreatment with CoQ10 (30 mg CoQ10/kg diet) together with *L*-carnitine (2.8 mg carnitine/kg diet), alpha-tocopherol (30 IU vitamin E/kg diet) and selenium (1 mg selenium as sodium selenite/kg diet), decreased DNA damage and the activities of Ca^{2+} , aspartate aminotransferase (ASAT) and alanine aminotransferase (ALAT) in the liver. However, the CoQ10 alone did not significantly protect against toxic cell death and glutathione depletion induced by FB_1 (Atroshi et al. [1999\)](#page-17-21). Another

study indicated that S-adenosylmethionine (SAM) and methylthioadenosine (MTA) were ineffective in protecting against $FB₁$ hepatotoxicity in C57BL/6 N mice. Neither SAM (intraperitoneal injection of 25 mg/kg b.w.) nor MTA (intraperitoneal injection of 25 mg/kg b.w.) modified the FB_1 -induced expression of TNF α , IL-1 α or IL-1 receptor antagonist. Both agents prevented an increase in free sphingosine but not sphinganine, and they could not improve the GSH level in liver following $FB₁$ treatment (subcutaneous injection of 2.25 mg/kg b.w.) (He et al. [2006\)](#page-18-21).

Metabolism of FBs

Metabolic pathways

The metabolism of FBs has already been studied since the 1990s, and some metabolites have been identified in the past years. The metabolism of FBs can notably occur in the liver and the digestive tract. The major concerned metabolic pathways of FBs are hydrolysis, acylation and transamination (Fig. [4](#page-14-0)) (Hartinger et al. [2011;](#page-18-25) Humpf et al. [1998](#page-18-26); Seiferlein et al. [2007\)](#page-19-26).

 $HFB₁$ (also called AP₁), the major metabolite of FB₁, originates from $FB₁$ by the hydrolysis of tricarballylic acid side chains at carbon 14 and 15, which are replaced by hydroxyl groups (Hartinger et al. 2011). HFB₁ (42-121 µg/ kg) was detected in the liver of weaned piglets fed a diet with $FB₁$ at 30 mg/kg feed for 6 weeks based on high-performance liquid chromatography (HPLC) combined with fluorescence detection (FL), indicating that the appearance of $HFB₁$ in the liver clearly originates from the breakdown of $FB₁$ from dietary origin (Pagliuca et al. [2005](#page-19-27)). Humpf et al. ([1998\)](#page-18-26) reported that HFB_1 (but not FB_1 , under the conditions tested) can be acylated to N -palmitoyl-HFB₁ (also called *N*-Pal-HFB₁ or pHFB₁) with a V_{max}/km of approximately 1 by ceramide synthase (Humpf et al. [1998](#page-18-26)). As the hydrolyzed forms of FB_1 and FB_2 , HFB₁ and HFB₂ (also called AP_2), respectively, can be metabolized by ceramide synthase to their respective *N*-acylated metabolites (nervonoyl- or palmitoyl-HFB₁ and -HFB₂) using rat liver microsomes and palmitoyl-CoA or nervonoyl-CoA as cosubstrates; the formation of N -acyl-HFB₁ was further identified in vivo using a HPLC-MS/MS method to detect tissues from rats given intraperitoneal doses of $HFB₁$ (Seiferlein et al. [2007\)](#page-19-26). *N*-acyl- $FB₁$ derivatives were also detected as in cultured cells at levels of up to 10 pM/mg of protein, and the acyl-chain length of the N -acyl-FB₁ depends on the ceramide synthase isoform acylating them (Harrer et al. [2013](#page-18-27)). A recent study found that the *N*-acyl chain length of the $FB₁$ metabolites varied in a tissuedependent manner with C_{16} derivatives predominating in the kidney and C_{24} derivatives being prevalent in the liver

when male rats were dosed with intraperitoneal injection with 0.5, 1 or 2 mg/kg body weight $FB₁$ on five consecutive days (Harrer et al. [2015\)](#page-18-28).

Metabolizing enzymes

Encoded as part of a gene cluster of *Sphingopyxis* sp. MTA144, the aminotransferase FumI could degrade $FB₁$ and catalyze deamination of the first intermediate of the catabolic pathway, hydrolyzed $FB₁$. FumI showed activity in the range of pH 6–10 with an optimum at pH 8.5, and in the range of $6-50$ °C with an optimum at 35 °C, and at low salt concentration (Hartinger et al. [2011](#page-18-25)). Ceramide synthase also acylates HFB_1 to pHFB₁ with a V_{max}/km of approximately 1 (Humpf et al. [1998](#page-18-26)). Additionally, $HFB₁$ can be obtained by de-esterification of $FB₁$ with a carboxylesterase and the toxicity of $HFB₁$ was much lower than $FB₁$, indicating that conversion to $HFB₁$ could be a good strategy to reduce FB_1 exposure (Grenier et al. [2012\)](#page-17-27).

The metabolizing enzymes for $FB₁$ in ruminants were also investigated. However, it appeared that no metabolizing enzyme was involved in the metabolism of $FB₁$ because no biodegradation or metabolism of $FB₁$ by liver microsomes was detectable after incubating $FB₁$ with bovine microsomes in the presence of a regenerating system for 1 h (Spotti et al. [2001\)](#page-19-28). Additionally, an in vitro study showed that no hydrolyzed metabolites (aminopolyols or aminopentol) were detected when FB_1 (1 μ g/mL) was incubated in ruminal fluid for up to 72 h, in the presence or absence of alfalfa as a substrate for microbial growth, using a model rumen, suggesting that $FB₁$ is poorly metabolized in the rumen and such metabolism is not the cause of the tolerance to this toxin displayed by ruminants (Caloni et al. [2000](#page-17-28)).

Strong differences were noted concerning the effects of FB_1 on drug-metabolizing enzyme activities. FB_1 (0, 5, 15 and 45 mg/kg/day) can affect drug-metabolizing enzyme activities when four groups of five growing ducks were treated with daily oral administration over 12 days. It can significantly increase benzphetamine, ethylmorphine, erythromycin *N*-demethylases and ethoxyresorufin *O*-deethylase activities (respectively, by 114, 242, 57 and 27 % with 5 mg/kg/day and 1024, 969, 200 and 147 % with 45 mg/kg/day) except glutathione S-transferase (GST) and significantly increase aminopyrine and nitrosodimethylamine *N*-demethylases, methoxyresorufin and pentoxyresorufin *O*-dealkylases and UDP-glucuronyltransferase activities at the dose of 45 mg/kg/day; this suggests that phase 1 drug-metabolizing enzyme activities appeared more sensitive to the toxin than phase 2 (Raynal et al. [2001](#page-19-29)). When rats were given intraperitoneal administration at three dose levels (0.125, 0.25 and 2.5 mg/ kg) once a day for 6 days, $FB₁$ increased the renal and

Fig. 4 Metabolic pathways of fumonisins FB₁ and FB₂, effects of FB₁ on the metabolizing enzymes. Cited from (Harrer et al. [2013;](#page-18-27) Hartinger et al. [2011;](#page-18-25) Humpf et al. [1998](#page-18-26); Seiferlein et al. [2007](#page-19-26))

hepatic *N*-demethylation of erythromycin (CYP3A1) and the hepatic *O*-deethylation of ethoxyresorufin (CYP1A1) at the two highest doses and increased the renal *O*-deethylation of ethoxyresorufin at the highest dose of 2.5 mg/ kg. Furthermore, CYP4A subfamily, the beta-oxidation of palmitoyl-coenzyme A (CoA) and trans-2-enoyl-CoA hydratase were also increased in the treatment of $FB₁$ at the two highest doses in the liver (Fig. [4\)](#page-14-0) (Martinez-Larrañaga et al. [1996\)](#page-18-29).

Other factors on the metabolism of FBs

The cecal microbiota of pigs can also transform $FB₁$ to the metabolites. In an in vitro study, pure $FB₁$ was treated with the suspensions of cecal contents and McDougall buffer solution under anaerobic conditions for 0, 12, 24, 48 and 72 h. After 48 h, the conversion of $FB₁$ to partially hydrolyzed FB_1 (pHFB₁) (46 %) was nearly equal to the percentage ratio of FB_1 , while it was 49 % by 72 h. In vitro, the

conversion of FB_1 to HFB₁ was less than 1 %, indicating that the most probable metabolic process is the removal of one propane-1,2,3-tricarboxylic acid side chain by esterase action in the mammalian gut, while the conversion of $FB₁$ to HFB₁ is not a priority during metabolism (Fodor et al. [2007](#page-17-29)). Alkaline processing like nixtamalization of maize generates $pHFB_1$ and HFB_1 and thermal treatment in the presence of reducing sugars leads to the formation of *N*-(1-deoxy-p-fructos-1-yl) fumonisin₁ (NDF) (Grenier et al. [2012;](#page-17-27) Hahn et al. [2015](#page-17-30)). The investigated alkaline and thermal processing products of $FB₁$ were non-toxic for rats at the tested concentrations (10 mg/kg FB_1 , 7.8 mg/kg pHFB₁, 5.6 mg/kg HFB₁ and 12.2 mg/kg NDF), indicating that appropriate food processing can reduce fumonisin toxicity for humans (Hahn et al. [2015\)](#page-17-30).

The toxicity of FBs and their metabolites

 FB_1 , pHFB₁, pHFB₂ and HFB₁ metabolites presented poor toxicity for the human intestinal cell line Caco-2 when cells were treated with them (in the range of 1–138 mM) for 48 h (Caloni et al. 2002). Although FB₁ appears the most toxic to cells, $HFB₁$, losing the tricarballylic acid chain is more bioavailable than $FB₁$ on intestinal cells, indicating that risk evaluations of the metabolites of FBs are also necessary (Caloni et al. [2002](#page-17-31)). In another study, a significant increase in sphinganine (up to 7000 % compared to control cells) was observed with FB_1 , FB_2 , FB_3 , HFB_1 and $pHFB_1$, and sphingosine levels remained almost unchanged, indicating that all substrates effectively inhibited ceramide synthase. Furthermore, only exposure to 10 μ M/L FB₁ for 24 h led to a significant increase in caspase-3 activity, chromatin condensation and DNA fragmentation, suggesting that $FB₂$, $FB₃$ and the metabolites of $FB₁$ could not induce apoptosis of human proximal tubule-derived cells (IHKE cells), whereas FB_1 could (Seefelder et al. [2003\)](#page-19-30). A previous study revealed that $HFB₁$ showed much lower toxicity than $FB₁$. When piglets were exposed by gavage for 2 weeks to 2.8 μ M FB₁ or HFB₁/kg body weight/day, FB₁ induced significant toxicity as well as hepatotoxicity and the toxic effects on the morphology of the different segments of the small intestine, decrease in villi height and modification of intestinal cytokine expression, whereas $HFB₁$ did not trigger hepatotoxicity, did not impair intestinal morphology and slightly modified the intestinal immune response (Grenier et al. [2012\)](#page-17-27). However, another in vivo study, where rats received diets supplemented with $FB₁$ or $HFB₁$, showed that the latter can be more toxic than the parent compound and induces fumonisin-like liver and kidney lesions (Voss et al. [1996\)](#page-19-31). Furthermore, $HFB₁$ shows cancer-promoting activity in liver (Hendrich et al. [1993\)](#page-18-30). The different toxicity induced by $FB₁$ in piglets and rats suggested that there are species differences in the toxicity of $FB₁$, which was far from clear and the reason was worthy of further investigation.

 $p_{HFB₁}$ was found to be at least ten times more toxic than $FB₁$ or $HFB₁$ and caused sphinganine accumulation as an inhibitor of ceramide synthase using HT29 cells, showing that $pHFB_1$ was a toxic metabolite and may play a role in the diseases caused by FBs (Humpf et al. [1998](#page-18-26)). To investigate the toxic effect on sphingolipid metabolism, high purity FB_1 , pHFB₁a + b, HFB₁ and NDF were fed to male SD rats for 3 weeks, and the results showed that the Sa/So ratios were significantly increased in the $FB₁$ positive control group, and the Sa/So ratios of $pHFB_1$, HFB_1 and NDF were indifferent from the negative control group (Hahn et al. 2015). Considering the fact that 2-keto-HFB₁ cannot be acylated in a similar way to $HFB₁$ (Humpf et al. [1998](#page-18-26); Seiferlein et al. 2007), 2-keto-HFB₁ was speculated to be non-toxic (Hartinger et al. [2011\)](#page-18-25). The acylation products of HFB₁ (C₁₆-HFB₁ and C_{24:1}-HFB₁) and HFB₂ (C₁₆-HFB₂) and $C_{24:1}$ -HFB₂) also inhibited ceramide synthase and significantly reduced the number of viable cells in an in vitro MTT assay using HT29 cells, suggesting that acylation apparently does not detoxify hydrolyzed fumonisins (Seiferlein et al. [2007](#page-19-26)). A recent study confirmed that acylation of $FB₁$ was even more toxic than the parent $FB₁$ when comparing the cytotoxicity of 20 μ M *N*-acyl-FB₁ derivatives with that of $FB₁$ at the same concentration for 8 h in human embryonic kidney (Hek) cells, Hep3B cells and human skin fibroblasts, respectively (Harrer et al. [2013](#page-18-27)).

In summary, FBs was metabolized mainly through hydrolysis, acylation and transamination progress. The metabolites of FBs were different in species. Differences in regulated enzymatic profiles among the analyzed species may be the reasons for the different metabolism of FBs. Enzymatic detoxification of $FB₁$ may be a suitable approach with which to increase feed and food safety, because the main metabolites of $FB₁$ showed lower toxicity than $FB₁$. Furthermore, the cooking progress was also a good strategy to reduce FBs exposure.

Conclusion

FBs were considered to be one of the most important contaminants of corn and corn-based food worldwide. The most potent and abundant of the class of FBs is $FB₁$. FBs could cause a variety of species-specific toxicological effects in human, domestic and laboratory animals. Therefore, it was necessary to investigate the toxic effect of FBs and therefore to protect from the injury of these toxins.

Most of the studies regarding oxidative stress induced by FBs (mainly $FB₁$) have been carried out on in vitro models and in vivo studies. A recent study found that GSH level was increased in a dose-dependent manner after 24-h

incubation of astrocytes with $FB₁$. Furthermore, in experiments in which the pre-incubation of cells with a conjugator of GSH did not affect ROS production induced by $FB₁$, this indicated that ROS is not involved in $FB₁$ toxicity, at least in astrocytes (Domijan and Abramov [2011\)](#page-17-14). Similar conclusions were drawn by the previous studies (Abel and Gelderblom [1998;](#page-17-16) Galvano et al. [2002](#page-17-15)). However, no similar study has been carried out in other cells. However, in some cells, such as SMC, SH-SY5Y cells, U-118MG cells, mouse C6 glioblastoma cells and rat GT1-7 hypothalamic cells, ROS production was suggested to play important roles in the toxic effects of FB_1 ; it was not clear whether the ROS over-production was a consequence rather than a mechanism of toxicity. Moreover, discrepancies were noted in ROS generation or not in SH-SY5Y cells by Domijan and Abramov (2011) (2011) and by Stockmann-Juvala et al. [\(2004a,](#page-19-14) [b](#page-19-15)), suggesting that there are a few studies that should be carried out to determine the role of ROS in the toxic mechanism of $FB₁$.

The mechanism of toxicity of FBs is complex, but the major biochemical effect of FBs is the inhibition of ceramide synthase (sphinganine- and sphingosine-*N*-acyl transferase), a key enzyme in the sphingolipid metabolism (Seefelder et al. [2003](#page-19-30)). ROS and oxidative stress played important roles in cellular apoptosis. However, few studies were carried out to elucidate the role of the inhibition of ceramide synthase induced by $FB₁$ in the oxidative stress and ROS generation in vitro and in vivo. The inhibition of ceramide synthase is sufficient for the induction of apoptosis only in some cell types (Seefelder et al. [2003](#page-19-30)). Then, the factors playing a critical role in the different induction of cellular apoptosis induced by $FB₁$ were worthy of investigation. Moreover, the research into the relationship between oxidative stress, ROS production and other factors also contribute to reveal the toxic mechanism of $FB₁$.

In a study to evaluate the toxic effects of $FB₁$ and other FBs, including FB₂, FB₃, fumonisin P1, HFB₁, *N*-acetyl- $FB₁$ and *N*-carboxymethyl- $FB₁$ (approximately 0, 14, 70) and 140 μ M/kg diet) in mice for 28 days, only FB₁ showed hepatotoxic effects (Howard et al. 2002). However, FB₂ could cause equine leukoencephalomalacia (Ross et al. [1994](#page-19-32)). Furthermore, short-term carcinogenesis studies in the rat liver bioassay indicated that FB_2 and FB_3 closely mimic the toxicological and cancer-initiating activity of $FB₁$ and thus could contribute to the toxicological effects of FBs in animals (Gelderblom et al. [1992](#page-17-32)). Therefore, the toxic effects of other FBs, such as $FB₂$ and $FB₃$, in vitro and in vivo seemed to be worthy of evaluation. However, few studies were carried out to reveal the potentially toxic effects of other FBs besides $FB₁$.

Various studies, including in vitro models and in vivo studies, have identified that the induction of an oxidative stress response induced by FBs exposure was

considered one of the mechanisms responsible for its toxicity (Tables [1,](#page-4-0) [2\)](#page-6-0). However, most in vivo studies were carried on rats or mice, and only one study was performed in chicks. Thus, greater emphasis should be placed on in vivo studies of livestock and poultry with the doses, which are commonly encountered in food crops or in feed.

The in vitro effects of hydrolyzed FBs were reported to range from less to more toxic than those of the intact FBs. The *N*-acyl-metabolites are even more cytotoxic than $FB₁$, suggesting that the 2-amino group of hydrolyzed FBs might play a critical role for toxicity. As for the metabolism of FBs in animals, the metabolism of $FB₁$ in cattle was significantly different from that in piglets. In the rumen, some studies found that $FB₁$ was apparently not dependent on its detoxification in the liver, and it still remains unclear why the tolerance of ruminants to $FB₁$ was higher than that of pigs. $HFB₁$ was found to show much lower toxicity than $FB₁$ in piglets, whereas it was more toxic in rats than the parent compound and induced fumonisin-like liver and kidney lesions and showed cancer-promoting activity in the liver. It was necessary to compare the different metabolizing progress among the rumen of piglets and rats, and even more species. However, until now, the metabolizing difference in species was far from clear.

It was interesting to note that oxidative stress occurred in rats, mice and chicks in the toxicology studies of $FB₁$, and various compounds, including free radical scavengers, can efficiently combat FB-induced damage, suggesting that the toxic effects of $FB₁$ might be closely related to its metabolizing progress and oxidative stress. The main metabolizing route of FBs was hydrolysis, acylation and transamination. However, it was far from clear which kind of metabolizing progress was closely related to oxidative stress, which was worthy of further investigation. Understanding the relationship between the metabolism of FBs and oxidative stress and identifying the related enzymes will contribute more to combating the toxicity induced by FBs.

To protect cells from FB_1 -induced oxidative stress, a variety of compounds have been evaluated for their antioxidative effects. Further understanding the role of oxidative stress as well as the metabolism of FBs in FB-induced toxicity will throw new light onto the use of antioxidants and scavengers of ROS. Considering the worldwide containment of FBs, the work focused on looking for highly effective antioxidants and efficient detoxification enzymes seems to be very meaningful to reduce the various toxic effects induced by FBs.

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Compliance with ethical standards

Conflict of interest The authors declare that there are no conflicts of interest.

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