

Fumonisin B1 + B2 change the expression of genes in apoptosis balance in Nile tilapia fingerlings

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ABSTRACT

In this study, we tested the hypothesis that the ingestion of feed containing fumonisin affects the apoptosis balance in liver cells. Thus, a trial was conducted with Nile tilapia fingerlings (2.64 ± 0.06 g) fed diets containing increasing levels of fumonisin (FB) (0-control diet, 20, 40, and 60 mg/kg), and expression of caspase7 (CASP7) and sphingosine phosphate lyase (SPL) genes in liver tissue was evaluated at 15 and 30 days of treatment. SPL:CASP7 mRNA ratio was also evaluated. Relative mRNA levels of SPL decreased in tilapia treated with 60 mg FB/kg of diet at 15 days ($P < .0001$), while a significant increase in SPL was detected at the two highest levels of inclusion at 30 days ($P < .0001$). CASP7 gene expression decreased linearly as fumonisin was added, at 15 days ($P < .0001$). At 30 days, however, the relative CASP7 mRNA levels were only reduced in the tilapia treated with 60 mg FB/kg of diet ($P = .0011$). The results obtained in this study suggest a direct relationship between SPL and CASP7 expressions, so it can be inferred that as the period of consumption and level of fumonisin are increased, the numerical relationship between SPL:CASP7 mRNA is also increased.

1. Introduction

Fumonisin is a mycotoxin produced during the secondary metabolism of the fungi *Fusarium verticillioides* and *F. proliferatum* (Turner et al., 1999). The presence of fumonisin in animal feeds is relatively common, since corn, that should be a natural target of contamination by the mycotoxins, is commonly used as an energy source (Griessler and Encarnação, 2009). It is known that different levels of fumonisin in fish may affect weight gain (Gbore et al., 2010; Tuan et al., 2003; Pepeljnjak et al., 2002; Lumlerdacha et al., 1995), cause hematological disorders (Gbore et al., 2010; Pepeljnjak et al., 2002), neurotoxicity (Kovačić et al., 2009), and change the relative concentrations of sphinganine (Sa) and sphingosine (So) (Goel et al., 1994).

The most widely accepted theory fumonisin's action is that this mycotoxin competes with sphingoid bases Sa and So for the binding site on the ceramide synthase enzyme, resulting in reduction of relative concentrations of ceramide, an increase in the level of free sphingoid

bases (Sa and So) and subsequently of sphingosine 1-phosphate (S1P) (Yoo et al., 1992; Wang et al., 1991).

The imbalance between these bases has been identified as the main factor associated with the hepatotoxic effects of fumonisin, and may be associated with changes in levels of caspases, enzymes involved in triggering the cascade of apoptotic events. The role of fumonisins in apoptotic balance has been reported as being controversial — at times they have been identified as pro-apoptotic (Ribeiro et al., 2010; Jones et al., 2001), and at other times as anti-apoptotic (Boppana et al., 2014; Mullen et al., 2012; Bose et al., 1995). In some cases, fumonisins have been identified as inducers of cell-death resistance even after cells have been subjected to stress factors that represent a risk to the organism (Mullen et al., 2011; Biswal et al., 2000).

Levels of ceramide, sphingoid bases, and S1P are directly associated with cell proliferation and apoptosis. Excess Sa and So can be minimized through their bioconversion to S1P (Riley and Voss, 2006). This bioconversion may disrupt the apoptosis balance by generating large

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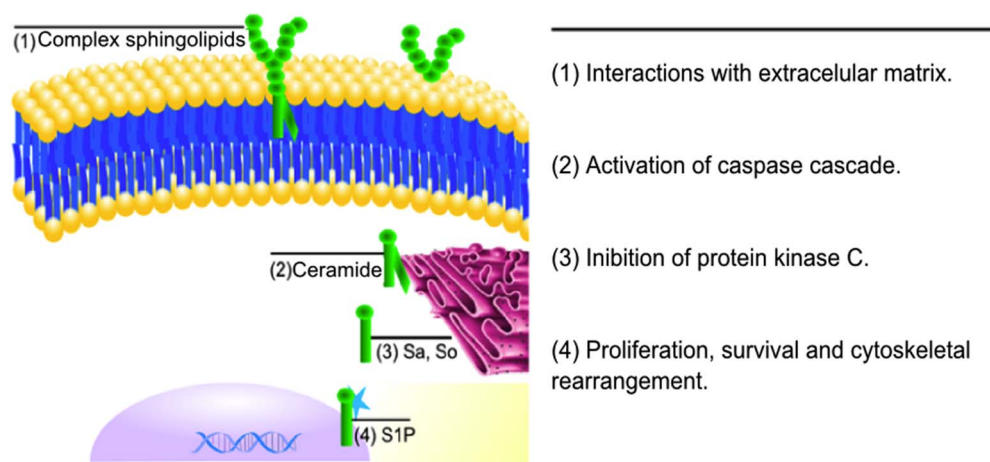


Fig. 1. Participation of complex sphingolipids and intermediates in processes related to cell proliferation. Adapted from (Spiegel and Merrill Jr, 1996).

concentrations of S1P, as this compound is known to be a bioactive metabolite capable of promoting cell proliferation and survival by inhibition of apoptosis (Taha et al., 2006) (Fig. 1).

Therefore, it is likely that fumonisins cause a reduction of ceramide, accumulation of sphinganine, and elevated production of S1P, thereby changing the apoptosis balance. Under normal conditions, homeostasis between sphingosine, sphinganine, and S1P is strictly regulated by activity of several key enzymes, e.g., sphingosine kinase (SPK), sphingosine-1-phosphate phosphatase (S1PP), and sphingosine phosphate lyase (SPL). Sphingosine is phosphorylated by SPK to form S1P. In this stage, S1P may be reverted to sphingosine by S1PP, or even irreversibly generate phosphoethanolamine and hexadecenal by means of SPL action (Taha et al., 2006). The function of phosphoethanolamine is not well known yet; nevertheless, it is believed that the products generated by SPL may be related to an increase in apoptosis by a caspase-3-independent signaling pathway (Ferreira et al., 2013).

In fish, there are no literature reports of fumonisin's action on the molecular mechanisms involved in apoptosis balance such as caspase and SPL gene expression. In this study, we tested the hypothesis that ingestion of feed containing increasing levels of fumonisin mycotoxin affects expression of genes related to apoptosis balance according to period of consumption. Thus, a trial was conducted with Nile tilapia fingerlings fed diets containing different levels of fumonisin to evaluate expression of caspase 7 (CASP7) and sphingosine phosphate lyase (SPL) genes in liver tissue at 15 and 30 days of treatment.

2. Materials and methods

2.1. Ethics statement

Study protocols were approved by Ethics Committee on Animal Use (CEUA - UniCesumar), Maringá, PR - Brazil. Fishes were anesthetized by immersion in eugenol solution (350 μ L/L) prior to sampling and livers were extracted based on the Guide to Care and Use of Laboratory Animals in Brazil.

2.2. Experimental design and animals

The experiment was conducted in the laboratory of Aquaculture of the State University of Maringá (UEM), located in Maringá-PR, Brazil, during the entire month of October. A total of 180 sex-reversed male fingerlings of GIFT \times Thai breed group, with an average weight of 2.64 ± 0.06 g, were used in the experiment. Fishes were distributed into three fiber cement tanks with a useful volume of 870 L each, an individual water restoration system (15%/day) and constant aeration provided through porous rock coupled with a central air blower. Four hapa net cages were inserted into each tank, totaling 12 experimental

units with four treatments and three replicates. Each hapa net cage had an individual volume of 217.50 and housed 15 fishes, totaling 14.5 L/fish. Fishes went through an acclimation period of 15 days before the beginning of the experiment.

2.3. Water quality parameters

In each cage, temperature was measured twice a day, at 09 h00 and 17 h00. Dissolved oxygen and pH were measured in the morning during the entire experimental period using an individual colorimetric kit.

2.4. Experimental diets

Four experimental groups were formed: GROUP 1 - control diet, with 0.0 mg of inclusion of FB/kg of diet; GROUP 2–20 mg FB/kg (16.36 mg FB₁ + 3.64 mg FB₂); GROUP 3–40 mg FB/kg (32.71 mg FB₁ + 7.29 mg FB₂); and GROUP 4–60 mg of inclusion of FB/kg (49.07 mg FB₁ + 10.93 mg FB₂).

Four isocaloric (approximately 3000 kcal of digestible energy/kg of diet) and isoproteic (approximately 33% crude protein) diets were formulated, varying only in inclusion of different levels of fumonisin (B1 + B2) (FB₁ - 6.06 mg toxin/g of medium and FB₂ - 1.35 mg toxin/g of medium, totaling a concentration of 7.41 mg toxin/g of medium) (Table 1). Growth medium was obtained from and evaluated in the Laboratory of Mycotoxicological Analysis (LAMIC) at the Federal University of Santa Maria (UFMS), based on *F. verticillioides* fungus, strain MRC 826.

Feed was pelletized, dried in a forced-air oven at 55 °C for 48 h, and then crumbled in a manual mill. Particles were classified according to their size (1 to 2 mm) and concentrations of fumonisin were later confirmed by laboratory analysis (LAMIC). Diets were hand-fed three times a day until apparent satiety of fish.

2.5. Gene expression

For analysis of gene expression, livers of six animals from each treatment were collected at 15 and 30 days of experiment to “RNA holder” tubes (BioAgencyBiotecnologia, São Paulo, Brazil) and stored at –80 °C until total RNA extraction.

Total RNA was extracted using Trizol® (Invitrogen, Carlsbad CA, USA), according to manufacturer's instructions. All materials used for extraction were pretreated with RNase AWAY® (Invitrogen, Carlsbad, CA, USA), an RNase inhibitor. Tissue was weighed, ground, and mixed with Trizol (1 mL of Trizol per 35 mg of tissue) until it was completely dissociated. After this step, 200 mL of chloroform was added to the samples, and the mixture was manually homogenized for 1 min. Next, samples were centrifuged for 15 min at 12,000 rpm and 4 °C, forming

Table 1
Composition of the experimental diets.

Ingredient (%)	Experimental diet			
	Control	20 mg FB/kg	40 mg FB/kg	60 mg FB/kg
Soybean	53.7	53.7	53.7	53.7
Corn	31.0	31.0	31.0	31.0
Corn gluten 60	8.2	8.2	8.2	8.2
Soybean oil	2.0	2.0	2.0	2.0
Corn starch	1.0	1.0	1.0	1.0
DL-Methionine	0.01	0.01	0.01	0.01
Calcitic limestone	0.1	0.1	0.1	0.1
Dicalcium phosphate	2.9	2.9	2.9	2.9
Vitamin C ^a	0.06	0.06	0.06	0.06
Mineral-vitamin supplement ^b	0.5	0.5	0.5	0.5
Common salt	0.5	0.5	0.5	0.5
BHT (antioxidant) ^c	0.03	0.03	0.03	0.03
Fumonisin B1 + B2	–	0.002	0.004	0.006
TOTAL	100	100	100	100
Chemical composition of the diet (%) ^d				
Crude protein	33.62	33.60	33.61	33.61
Dry matter	89.52	89.51	89.50	89.52
Fat	4.33	4.33	4.30	4.29
Ashes	5.79	5.75	5.76	5.78

^a Vitamin C: calcium salt L-ascorbic acid 2-phosphate, 42% active ingredient.

^b Mineral-vitamin supplement: Composition/kg of product: vit. A = 1200,000 IU; vit. D3 = 200,000 IU; vit. E = 12,000 mg; vit. K3 = 2400 mg; vit. B1 = 4.800 mg; vit. B2 = 4800 mg; vit. B6 = 4000 mg; vit. B12 = 4800 mg; folic acid = 1200 mg; calcium pantothenate = 12,000 mg; vit. C = 48,000 mg; biotin = 48 mg; choline = 65,000 mg; nicotinic acid = 24,000 mg; Fe = 10,000 mg; Cu = 600 mg; Mn = 4000 mg; Zn = 6000 mg; I = 20 mg; Co = 2 mg; and Se = 20 mg.

^c Butylated hydroxytoluene (Antioxidant).

^d Composition on a fresh matter basis.

separate layers. During this stage, 500 µL of aqueous phase were collected and transferred to a clean tube containing 500 µL of isopropanol. Tubes were homogenized and centrifuged for 15 min at 12,000 rpm and 4 °C. Supernatant was discarded, while 1 mL of 75% ethanol was added to the precipitate for cleaning. Tubes were centrifuged once again at 12,000 rpm for 5 min, and supernatant was discarded. Pellet was then dried for 15 min at 4 °C and resuspended in RNase-free ultrapure water.

Total RNA concentration was measured in a spectrophotometer at 260 nm wavelength. RNA integrity was evaluated using agarose gel stained 1.5% with ethidium bromide (10%) and visualized under ultraviolet light. RNA samples were treated with DNase I (Invitrogen, Carlsbad, CA, USA), in accordance with the instructions of manufacturer to remove possible contaminations with genomic DNA.

SuperScript™ III First-Strand Synthesis Super Mix kit (Invitrogen Corporation, Brazil) was used for cDNA synthesis according to the manufacturer's instructions. In this stage, 6 µL of total RNA, 1 µL of oligo dT (50 µM oligo (dT) 20), and 1 µL of annealing buffer were added to a RNA-free sterile tube. Tubes were incubated for 5 min at 65 °C and immediately placed on ice for 1 min. Next, 10 µL 2× First-Strand Reaction Mix and 2 µL of a solution containing SuperScript™ III reverse transcriptase enzyme were added to the tubes. The resulting solution was incubated for 50 min at 50 °C and for 5 min at 85 °C, then immediately placed on ice. After this stage, samples were stored at –20 °C until their use.

Real-time PCR analyses were performed using the SYBR Green fluorescent dye (SYBR® GREEN PCR Master Mix, Applied Biosystems, USA).

Primers utilized for evaluation of caspase 7 (CASP7) and sphingosine phosphate lyase (SPL) genes were designed based on the sequences of genes stored at www.ncbi.nlm.nih.gov (access numbers XM_005471473 and XM_003441226.2, respectively), available through the website www.idtdna.com. Three endogenous controls obtained from Yang et al. (2013) were tested: UBCE (ubiquitin-conjugating

Table 2
Sequence of primers used for the real-time PCR reactions.

Primer	Sequence (5'-3')	Amplicon (bp)
CASP7 – F	GATTCCTGTGTGGTCTCTATG	94
CASP7 – R	GGTCAGTCTGTGGCATTATTA	
SPL F	GTCACCTGACCAAACCTCTAGTG	125
SPL R	GACAACCTCTGCTCCATCTT	
EF1-α – F	GCACGCTCTGCTGGCCTTT	250
EF1-α – R	GGCTCAATCTTCCATCCC	
UBCE – F	CTCTCAAATCAATGCCACTTCC	130
UBCE – R	CCCTGGTGGAGGTTCTTGT	
BACT – F	TGGTGGGTATGGGTCAGAAAG	217
BACT – R	TGTTGGCTTTGGGGTTCA	

enzyme - XM_003460024), EF1A (elongation factor1α - AB075952), and ACTB (β actin - XM_003455949). Of these, ACTB was selected, as it showed the best efficiency (between 90 and 110%) and did not vary statistically between the treatments. Primers utilized in this experiment are shown in Table 2.

All reactions were performed in the final volume of 12.5 µL, as well as in duplicates. Analysis of dissociation curves did not reveal any presence of primer dimers or non-specific products.

2.6. Statistical analysis

To evaluate relative expression, real-time PCR analysis data were transformed using $2^{-\Delta\Delta CT}$, as demonstrated by Livak and Schmittgen (2001). Gene expression data were evaluated using a completely randomized design in a 2 × 4 factorial arrangement (two ages and four levels of fumonisin inclusion), employing GLM procedure, and mean values were compared by Tukey's test ($P < .05$) (SAS Inst. Inc., Cary, NC, USA). Results were expressed as means and standard deviations. Data on ratio between mRNA expressions of SPL and CASP7 were analyzed using REG statement (SAS Inst. Inc., Cary, NC, USA).

3. Results

Mean values and standard deviations for relative expression of CASP7 and SPL genes in liver of Nile tilapia fingerlings according to level of inclusion of fumonisin and period of consumption are shown in Table 3.

Gene expression of SPL enzyme was significantly changed by diets. An interaction effect between period of consumption and levels of fumonisin in diet (Fig. 2) was detected. Relative SPL mRNA levels were reduced only in tilapia treated with the 60 mg FB/kg diet at 15 days ($P < .0001$), whereas at 30 days, SPL had a significant increase in case of the two highest levels of inclusion ($P < .0001$).

Gene expression for CASP7 was also significantly changed by diets, with the presence of a significant effect between period of consumption and levels of fumonisin (Fig. 3). At 15 days of treatment, inclusion of increasing levels of fumonisin reduced relative values of CASP7 mRNA in a linear manner ($P < .0001$). At 30 days, the relative CASP7 mRNA levels were only reduced in tilapia treated with the 60 mg FB/kg diet ($P = .0011$).

A greater expression of CASP7 mRNA as compared with SPL was observed for control treatment both at 15 and 30 days. Paradoxically, SPL mRNA levels were higher than those of CASP7 for the two highest levels of fumonisin tested, as demonstrated in Fig. 4.

When the numerical relationship between relative expressions of SPL and CASP7 was evaluated, a strong upward trend was observed for this ratio as a function of the period and levels of fumonisin inclusion (Fig. 5).

Table 3
Relative gene expression of the CASP7 and SPL genes at 15 and 30 days of ingestion of increasing levels of fumonisin.

Treatment	CASP7 (mRNA) AU ^a		SPL (mRNA) AU	
	15 days	30 days	15 days	30 days
Control	0.023 ± 0.01	0.0025 ± 0.0004	0.0051 ± 0.0003	0.0015 ± 0.0003
20 mg FB/kg	0.011 ± 0.0023	0.0024 ± 0.0006	0.0045 ± 0.0006	0.0013 ± 0.0001
40 mg FB/kg	0.004 ± 0.0014	0.0022 ± 0.0001	0.0044 ± 0.0009	0.0034 ± 0.0010
60 mg FB/kg	0.001 ± 0.0005	0.0007 ± 0.0001	0.0023 ± 0.0003	0.0023 ± 0.0006

^a AU - arbitrary units.

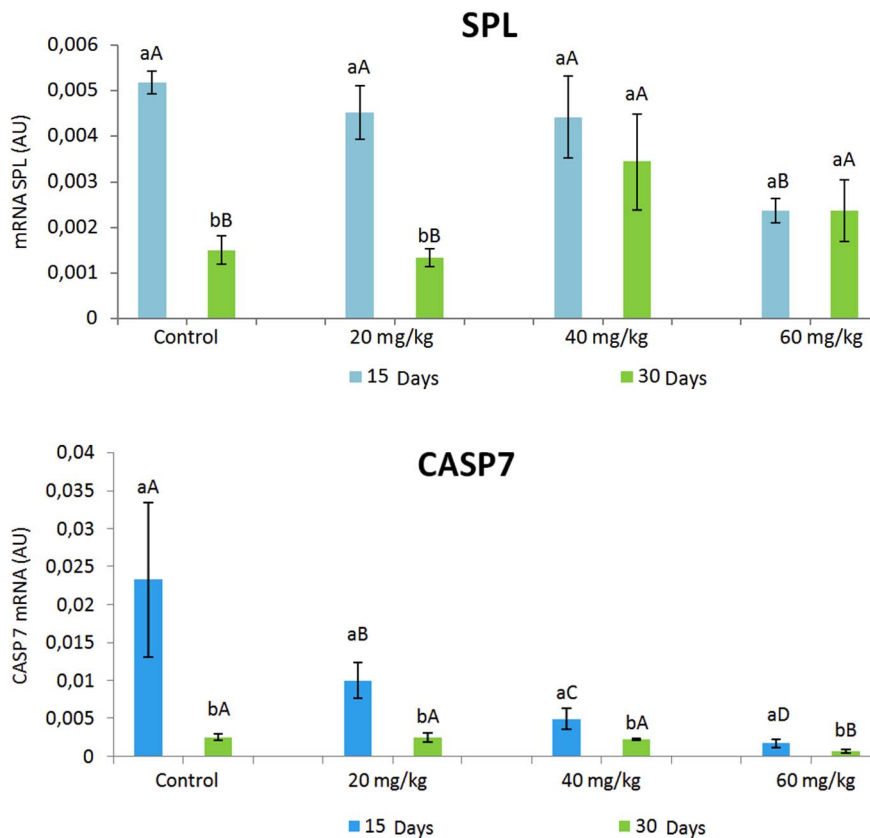


Fig. 2. Effects of period of consumption and levels of inclusion of fumonisin on the gene expression of SPL. Values are expressed in arbitrary units (AU). Lowercase letters on the bars compare the means between periods in the same treatment. Uppercase letters on the bars represent the test of means between the levels in the same period. Tukey's test at 5% probability.

Fig. 3. Effects of period of consumption and levels of inclusion of fumonisin on the gene expression of CASP7. Values are expressed in arbitrary units (AU). Lowercase letters on the bars compare the means between periods in the same treatment. Uppercase letters on the bars represent the test of means between the levels in the same period. Tukey's test at 5% probability.

4. Discussion

In this study, we evaluated gene expression of SPL and caspase 7 (CASP7) enzymes. For SPL, an effect of interaction between concentrations of fumonisin and days of exposure to the toxin ($P < .0001$) was observed, suggesting that the effects of fumonisin are dose-time-dependent. At 15 days, different levels of fumonisin affected mRNA expression of SPL ($P = .0002$), with a slight reduction of this expression only for animals that consumed diet with the highest level of fumonisin inclusion. On the other hand, in animals fed for 30 days, a significant increase in expression of SPL was discovered at the two highest tested levels of fumonisin inclusion ($P < .0001$) (Fig. 2).

S1P levels in cells are strongly regulated by the balance between its synthesis and degradation. It is possible that the response of gene expression of SPL enzyme at 15 days results from the activation of compensatory mechanism, in which the Sa/So-1-phosphate phosphatase enzyme (S1PP), which catalyzes return of sphingosine 1-phosphate to sphingosine, prevented accumulation of S1P, especially at the highest level of fumonisin inclusion, in an unsuccessful attempt to form ceramide via ceramide synthase.

At 30 days, an increase in SPL expression was observed. A longer duration of fumonisin exposure likely affected ceramide synthase,

promoting accumulation of sphingoid bases and consequently an increase in the production of S1P and SPL expression, as suggested by Loetscher et al. (2013). Also, biochemical signals observed with consumption of fumonisin, like Sa/Sa ratio, concentration of ceramide, and concentration of macromolecules involved in this pathway are time- and dose-dependent, corroborating the differences observed at 15 and 30 days of testing (Wang et al., 1992). There is a threshold between cell behavior that leads to apoptosis or proliferation. Huang et al. (2011) suggested a model called “sphingolipid rheostat”, in which the ceramide/sphingosine and S1P levels, which have opposing functions, are interchangeable within cells and may lead to cell death (when the balance is shifted towards ceramide/sphingosine) or survival and proliferation (when S1P levels are increased) (Fig. 6).

Thus, it is possible that the increase in levels of SPL observed at 30 days is an attempt of the body to reduce cell proliferation rate through irreversible catalysis of S1P. It is believed that SPL has the ability to change cell balance by its direct action on S1P levels, involving p53 and p38MAPK signaling pathways, which are associated with the signaling of cell death involving the caspases (Giovannini et al., 2011).

Some studies have identified fumonisins as stimulators of apoptosis via caspase (Kim et al., 2007; Gopee and Sharma, 2004); nevertheless,

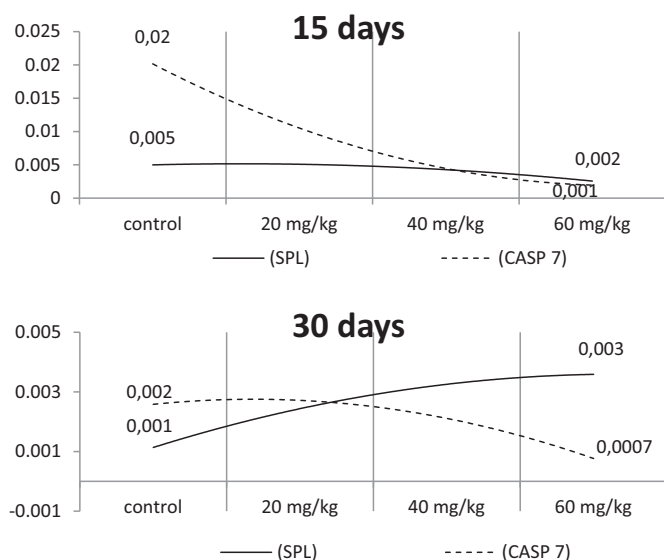


Fig. 4. Comparison between the expression pattern of SPL and CASP7 mRNA at 15 and 30 days of treatment.

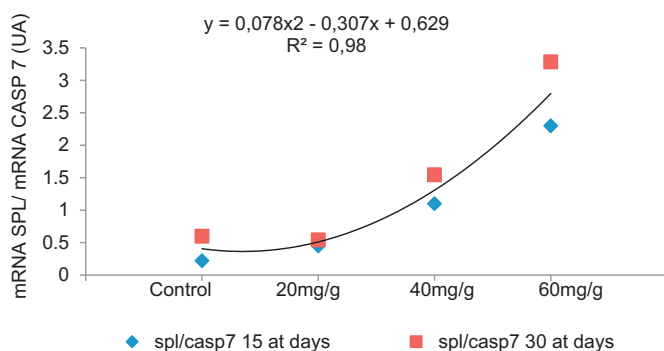


Fig. 5. Numerical relationship between the SPL and CASP7 mRNA levels in the liver of Nile tilapia challenged with increasing levels of fumonisin for 15 and 30 days.

the findings of this study suggest a different response. In the present experiment, fish subjected to increasing levels of fumonisin for 15 days showed a marked reduction of CASP7 mRNA (approximately 23 times). At 30 days of treatment, a reduction in CASP7 expression was also observed, but to a lesser extent (approximately 3.5 times) (Table 3). Overall, the resulting data demonstrates that inclusion of increasing levels of fumonisin reduces caspase 7 mRNA expression in liver of Nile tilapia. Similarly to our findings, Mullen et al. (2012) observed that inhibition of ceramide synthase by fumonisins also inhibited activation of caspases 3 and 7, with a subsequent decrease in apoptosis, suggesting that the roles of ceramide affect activation of caspase directly by influencing permeability of external mitochondrial membrane, preventing mobilization of factors responsible for the onset of the caspase cascade.

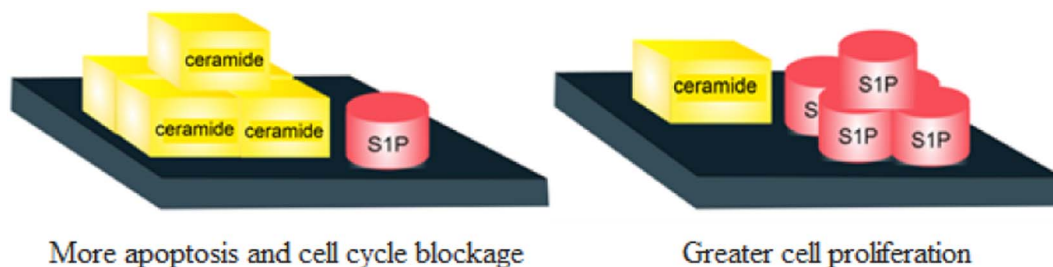


Fig. 6. Relationship between ceramide and S1P. Adapted from Huang et al. (2011).

In cells under normal conditions, there is an accumulation of intracellular ceramide after receiving genotoxic stimuli that acts directly on protein phosphatase 2 (PP2A) (Xin and Deng, 2006; Bionda et al., 2004). This phosphatase, together with anti-apoptotic Akt/PKB kinase, regulates the action of anti-apoptotic factors Bax/Bcl-2, promoting mitochondrial permeability (Taha et al., 2006). Subsequently, protein “Smac/DIABLO” and cytochrome c are released by the mitochondria, promoting activation of caspase cascade through the adapter protein Apaf-1 (Du et al., 2000). Boppana et al. (2014) demonstrated in their study that fumonisin plays an anti-apoptotic role by inhibiting accumulation of ceramide in endoplasmic reticulum, Bax translocation to mitochondria, and release of cytochrome c.

In this context, lower concentrations of ceramide resulting from consumption of fumonisin may be responsible for lower CASP7 expression observed in the present study and consequent reduction of caspase-dependent apoptosis rates. Our study suggests that fumonisin plays an anti-apoptotic role by means of expressive reduction of CASP7.

When SPL and CASP7 expressions for proposed levels of inclusion were evaluated synergistically, we could observe that a gene-expression pattern occurred, in which the control treatment resulted in higher CASP7 levels than SPL, while at the highest level of inclusion of fumonisin this response was the opposite, with a higher SPL expression than CASP7 (Fig. 4). Inversion of expression of these genes was more marked at 30 days. Kim et al. (2007) also observed an inverse pattern of expression for SPL and caspase 3 in liver of rats exposed to fumonisin; however, the lowest expression of caspase and the highest SPL expression were observed in animals subjected to control treatment.

In this study inverse patterns of expression for CASP7 and SPL were observed according to levels of fumonisin, so we also evaluated response shown by ratio between these genes. While analyzing this ratio, we observed an upward trend in this value as a response to increasing levels of fumonisin in diets (Fig. 5). Thus, it can be assumed that the mechanisms involved in increase of the SPL expression are also related to reduction of CASP7 expression, when animals consume a ceramide suppressant.

Overall, our results indicate alterations in SPL expression of tilapia fingerlings consuming increasing doses of fumonisin in diet and that the effects are dose- and time-dependent. It is likely that the observed reduction in CASP7 mRNA levels is dependent on the reduction in ceramide levels (Reiss et al., 2004), which were probably changed as a result of fumonisin action. The increase in SPL is probably the result of an increased number of sphingoid bases, and consequently of S1P. These results demonstrate a direct relationship between SPL and CASP7 expressions, so it can be inferred that as the time consuming fumonisin is increased, the numerical relationship between of SPL and CASP7 gene expressions will also increase.

5. Conclusion

Consumption of feed containing increasing levels of mycotoxin fumonisin B1 + B2 affects expression of genes involved in apoptotic balance, caspase 7 and sphingosine phosphate lyase, according to the period of ingestion.

Conflict of interest

The authors declare that they have no conflict of interest.

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