## Università degli Studi di Milano

PhD Course in Veterinary and Animal Science



# In vitro toxicological effects of Fumonisin B<sub>1</sub> alone and combined with other mycotoxins

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"I am ignorant of absolute truth. But I am humble before my ignorance and therein lies my honor and my reward."

Khalil Gibran

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## 1. Abstract

Mycotoxins, secondary metabolites produced by moulds, mainly *Aspergillus* spp., *Fusarium* spp. and *Penicillium* spp., are common contaminants of food and feed. The aim of this project was to evaluate: (i) the potential endocrine disruptor effects of fumonisin  $B_1$  (FB<sub>1</sub>), beauvericin (BEA), deoxynivalenol (DON) and zearalenone (ZEA) metabolites  $\alpha$ -zearalenol ( $\alpha$ -ZEA) and  $\beta$ -zearalenol ( $\beta$ -ZEA), alone and combined, using a bovine granulosa cell (GC) *in vitro* model and (ii) the individual and combined effects of FB<sub>1</sub> and BEA on the intestinal barrier using Caco-2 cells cultured *in vitro* on semipermeable inserts.

The results obtained indicated that FB<sub>1</sub> alone at all tested doses (0; 0.5; 1; 1.5; 3; 6  $\mu$ M) had no effects on GC proliferation and progesterone (P4) production. In the presence of  $\beta$ -ZEA at30 ng/mL (0.094  $\mu$ M), FB<sub>1</sub> at 30 ng/mL (0.042  $\mu$ M) showed a stimulatory effect on GC numbers. Cell proliferation decreased after exposure to  $\beta$ -ZEA alone at 5.0 mg/mL (15.6  $\mu$ M) and FB<sub>1</sub> with  $\alpha$ -ZEA and  $\beta$ -ZEA at the same concentration. Regarding steroid production, FB<sub>1</sub> at 30 ng/mL (0.042  $\mu$ M) and 100 ng/mL (0.13  $\mu$ M amplified the inhibitory effect of  $\beta$ -ZEA at 30 ng/mL (0.094  $\mu$ M) on estradiol (E2) production, while FB<sub>1</sub> alone increased (P<0.05) IGF1-induced E2 production. FB<sub>1</sub> in combination with  $\beta$ -ZEA decreased (P < 0.05) E2 production. FB<sub>1</sub> at 1, 1.5 and 3  $\mu$ M slightly inhibited (P < 0.05) E2 production. BEA at concentrations  $\geq$  3  $\mu$ M was found to strongly decrease (P < 0.05) both steroid production and FB<sub>1</sub> did not influence the effects of BEA. At 10  $\mu$ M both mycotoxins decreased (P < 0.001) serum-induced GC proliferation. At 30  $\mu$ M, BEA showed inhibitory effects on FSH plus IGF-1-induced CYP11A1 and CYP19A1 mRNA abundance (P < 0.05), whereas FB<sub>1</sub> at 30  $\mu$ M had no effect on CYP11A1 and CYP19A1 gene expression.

As regards the effects of FB<sub>1</sub> and BEA, alone and combined, on the Caco-2 intestinal barrier model data showed a TEER decrease after 1 h and 2 h of Bl exposure to BEA at 0.5 and 1.5  $\mu$ M and after 24 h of Bl exposure to BEA at 0.5  $\mu$ M, whereas after 24 h of Bl exposure, BEA at 3 and 6  $\mu$ M was found to significantly (P < 0.05) increase TEER.

FB<sub>1</sub> had no effect on the intestinal barrier integrity and when combined with BEA the TEER increase induced by BEA was no longer observed.

Cytokine release was observed only after exposure to BEA alone, and not in combination with FB<sub>1</sub>, with an increase of IL-6 and IL8 release after apical exposure to 3 and 6  $\mu$ M and after basolateral exposure to 1.5, 3, 6  $\mu$ M for IL-6 and only to 6  $\mu$ M for IL-8. TNF-  $\alpha$  release was induced by Ap (0.5 -1.5  $\mu$ M) and Bl (1.5  $\mu$ M) exposure to BEA.

Overall, these results provide information on *in vitro* toxicological effects of *Fusarium* mycotoxins.

## 2. General introduction

## 2.1 Mycotoxins

Mycotoxins are secondary metabolites produced by molds that contaminate food and feed worldwide (Schollenberger et al., 2007). 25% of the world's crop production is contaminated by mycotoxins and their global occurrence pose a serious risk to human and animal health (Pinton and Oswald, 2014). Mycotoxins produced by *Aspergillus, Fusarium*, and *Penicillium* genera are particularly significant for their ubiquity and toxicity (Osweiler, 2000).

There are several factors that influence the presence of mycotoxins in foods or feeds: extrinsic and intrinsic factors (Hussein and Brasel, 2001). Extrinsic factors include environmental conditions such as storage, temperature and humidity (Hussein and Brasel, 2001) while intrinsic ones include the strain specificity and the strain variation (Hussein and Brasel, 2001). Mycotoxins have a wide spectrum of toxic effects and the nature and the intensity of these effects are not only related to the time and dose of exposure, but also to the possible co-occurrence of mycotoxins that can result in additive, synergic or antagonistic toxicological effects (Fink-Gremmels, 1999). Chronic exposure to mycotoxins (Hussein and Brasel, 2001; Fink-Gremmels and Malekinejad, 2007) results in reduced feed intake, loss of body weight, immune-suppression with a subsequent increased susceptibility to infections, and possible effects on reproductive function (Fink-Gremmels and Malekinejad, 2007; Pestka, 2007; Voss et al., 2007).

## 2.2 Fusarium mycotoxins

Fusarium molds are commonly found on commodities in Europe, America and Asia and are considered the most prevalent species in the northern hemisphere (Creppy, 2002). Fusarium spp. synthesize a wide range of mycotoxins (Flannigan, 1991; Glenn, 2007) and the most important in terms of impact to animal health and production are fumonisins, trichothecenes and zearalenone (D'Mello et al., 1999; Uhlig et al., 2007; Jestoi, 2008).

Fusarium species require a moisture content of 25% (Newman and Raymond, 2005) and toxinogenesis is strongly influenced by this factor (Sweeney and Dobson, 1998). Consumption of a mycotoxin-contaminated diet may induce acute and chronic effects resulting in a broad variety of toxicological effects in animals (D'Mello et al., 1999; Binder et al., 2007; Smith, 2012; Cortinovis et al., 2014)

## 2.2.1 Worldwide contamination of Fusarium mycotoxins

The presence of mycotoxins depends on several factors such as their interactions with other organisms on the substrate where they coexist, the production site, humidity, temperature and the agricultural and post-harvest practices (Hussein and Brasel, 2001; Ferre, 2016). Several surveys reported the occurrence of mycotoxins relevant to the feed industry in different regions particularly with respect to fumonisins (Boutigny et al., 2012; Garrido et al., 2012) and deoxynivalenol (DON) (Boutigny et al., 2012; Grajewski et al., 2012). Rodrigues and Naehrer (2012) reported that fumonisins, DON and zearalenone (ZEA) were present in 64%, 59% and 45% of analyzed samples, respectively while, in a study conducted by Yoshinari et al. (2016) beuvericin (BEA) was found in 34% of the samples. As reported by Binder et al. (2007), more than half of the samples collected in Europe were contaminated as well as one third of the Asian-Pacific ones. The European Commission (EC) has established guidance levels for the presence of some Fusarium mycotoxins in animal feed (EC, 2006). For fumonisins (fumonisin B<sub>1</sub> and B<sub>2</sub>) the guidance values are 60 mg/kg for maize and maize products, 5 mg/kg for complementary and complete feedingstuffs for pigs, horses, rabbits and pet animals, 10 mg/kg for fish, 20 mg/kg for poultry, calves (<4 months), lambs and goat kids and 50 mg/kg for adult ruminants (>4 months) and mink (EC, 2006). Regarding DON the EC guidance values are 8 mg/kg for cereals and cereal products with the exception of corn by-products (12 mg/kg) and 5 mg/kg for complementary and complete feedingstuffs with the exception of feedingstuffs for pigs (0.9 mg/kg) and for calves (<4 months), lambs and goat kids (2 mg/kg) (EC, 2006; Pinton and Oswald, 2014). The ZEA EC guidance values recommended for complementary and complete feedingstuffs for piglets/gilts, sows/fattening pigs, and calves/dairy cattle/sheep/goats are 0.1, 0.25 and 0.5 mg/kg, respectively (EC, 2006; Streit et al., 2012). The ZEA EC guidance values for cereals/cereal products and corn by-products are 2 and 3 mg/kg, respectively (EC, 2006).

#### 2.2.2 Fumonisins

Fusarium proliferatum, Fusarium napiforme and Fusarium nygamai that have been shown to occur worldwide at significant levels in corn and corn by-products (Voss et al., 2007; Glenn, 2007). Fumonisins have been found to commonly occur in combination with deoxynivalenol (DON), zearalenone (ZEA) and beauvericin (BEA) in cereal grains and animal feed (Jestoi, 2008). Fumonisins are divided into four groups known as A, B, C and P (Marasas et al., 1984; Alberts et al., 1990; Yazar and Omurtag, 2008). The most

important is the B group that includes fumonisin B<sub>1</sub> (FB<sub>1</sub>), B<sub>2</sub> (FB<sub>2</sub>) and B<sub>3</sub> (FB<sub>3</sub>) with FB<sub>1</sub> (Fig. 2.1) being the most significant in terms of toxicity and occurrence (EFSA, different induces species-specific effects in animals leukoencephalomalacia in horses and pulmonary edema in pigs (Marasas et al., 1988; Ross et al., 1993; Voss et al., 2007). Cattle are considerably less sensitive to FB<sub>1</sub> than horses and pigs (Fink-Gremmels, 2008) and signs of liver and kidney injury have been reported in cattle only after exposure to very high concentrations of FB<sub>1</sub> (Osweiler et al., 1993; Baker and Rottinghaus, 1999; Mathur et al., 2001). In ruminants FB<sub>1</sub> has a very low bioavailability, is poorly degraded in the rumen (Caloni et al., 2000) and is found unmetabolized in feces (Cavret and Lacoeur, 2006). The mechanisms of toxicity for FB<sub>1</sub> are complex and may involve several molecular sites (Voss et al., 2007). FB<sub>1</sub> bears a remarkable structural resemblance to the long-chain (sphingoid) base backbones of sphingolipids and thus impairs sphingolipid biosynthesis (Wang et al., 1991; Merrill et al., 2001; Voss et al., 2002; Voss et al., 2007). The inhibition of ceramide synthase leads to the accumulation of sphingoid bases and to the depletion of complex sphingolipids, which interfere with the function of some membrane proteins (Wang et al., 1991; Merrill et al., 2001; Voss et al., 2002; Voss et al., 2007). Thus, the mechanism of toxicity of FB<sub>1</sub> is linked to its ability to disrupt the sphingolipid metabolism and the subsequent adverse effects on cell regulation (Wang et al., 1992; Marasas et al., 2004; Smith., 2012).

The International Agency for Research on Cancer (IARC) classified FB<sub>1</sub> as a possible carcinogen (class 2B) for humans (IARC, 2003).

**Figure 2.1:** Chemical structure of fumonisin B<sub>1</sub> (FB<sub>1</sub>).

#### 2.2.3 Trichothecenes

Trichothecenes are a family of *Fusarium* mycotoxins commonly found in cereal grains such as corn, wheat, rye, barley and oats. The family is subdivided into four groups (A, B, C and D) according to their chemical structure. Type A trichothecenes including T-2, HT-2 toxin, neosolaniol and diacetoxyscirpetol and type B including DON and its 3- acetyl and 15-acetyl derivates, nivalenol and fusarenon X are the most important (Escriva et al., 2015).

Trichothecenes inhibit the synthesis of RNA and DNA interfering with cellular metabolic activities and causing cell death (Thompson and Wannemacher, 1990) and are able to readily bind to eukaryotic ribosomes, in particular to the 60S ribosomal subunits, inhibiting protein synthesis (Pestka, 2010).

The general clinical signs of trichothecene toxicosis commonly include loss of appetite, vomiting, diarrhea, leukocytosis and gastrointestinal hemorrhage (Pestka and Smolinski, 2005; Maresca, 2013; Blajet-Kosicka et al., 2014; Pinton and Oswald, 2014). Ruminants are less sensitive to trichothecene toxicity whereas pigs and poultry are the most susceptible species to DON and T-2 toxin respectively (Pestka and Smolinski, 2005; Maresca, 2013; Pizzo et al., 2016).

## 2.2.4 Deoxynivalenol

Deoxynivalenol (DON), also known as vomitoxin, belongs to the type B group of trichothecenes and occurs commonly in grains such as maize and wheat but also in rice, oats and sorghum (Pinton and Oswald, 2014). DON has been implicated in farm animal disease outbreaks in many areas of the world and is considered one of the most hazardous food-associated mycotoxin for human health (Maresca, 2013; Pinton and Oswald, 2014).

Structurally, DON (Fig. 2.2) is a polar organic compound containing 3 free hydroxy groups (-OH), which are associated with its toxicity (Sobrova et al., 2010). The mechanism of action of DON is known to be related to its ability to bind eukaryotic ribosomes and inhibit protein synthesis by blocking translation and inhibiting the elongation of peptide chains (Larsen et al., 2004; Pestka, 2010). Moreover, DON can also induce cell apoptosis activating the mitogen-activated protein kinases (MAPKs) (Larsen et al., 2004; Pestka et al., 2010; Li et al., 2014). After ingestion of highly DON contaminated food common clinical signs include abdominal pain, vomit, diarrhea, leukocytosis and blood in stool. Loss of weight and altered immune function have been frequently observed after chronic exposure (Pestka and Smolinski, 2005).

DON was found to be able to cross the intestinal barrier modifying cellular functions and causing cell death in pigs (Maresca et al., 2013). *In vitro* studies carried out with the cell lines Caco-2, IPEC-1 and IPEC-J2 demonstrated that DON is able to impair the intestinal barrier function and may have serious consequences for human and animal health (Van de Walle et al., 2010; Awad et al., 2011; Vandenbroucke et al., 2011).

Ruminants are considered less susceptible to the adverse effects of DON because DON is rapidly converted in the rumen into de-epoxy DON (DOM1) that is less toxic compared to the parent compound (Cotè et al., 1986; Fink-Gremmels, 2008).

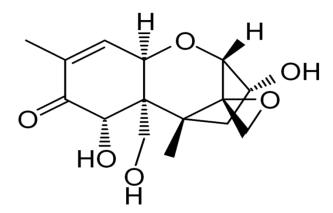


Figure 2.2: Chemical structure of deoxynivalenol (DON).

## 2.2.5 Zearalenone

Zearalenone (ZEA) (Fig. 2.3) is a widespread mycotoxin that is of great interest due to its toxic effects on animals (Fink-Gremmels, 1999; Fink-Gremmels and Malekinejad, 2007; Zinedine et al., 2007). ZEA is an estrogenic mycotoxin produced by several *Fusarium* species including *F. culmorum*, *F. graminearum*, *F. equiseti* and *F. crookwellense*. This mycotoxin commonly contaminates wheat, rice, maize, barley and other crops (Fink-Gremmels and Malekinejad, 2007). ZEA is a resorcyclic acid lactone, chemically described as 6-[10-hydroxy-6-oxo-trans-1-undecenyl]-B-resorcyclic acid lactone (Zinedine et al., 2007). This particular structure resembles several characteristics of steroid hormones and consents ZEA to bind estrogen receptor α (ESR1) and estrogen receptor β (ESR2) (Zinedine et al., 2007). In this way ZEA can act as an agonist and partial antagonist of estradiol (Malekinejad et al., 2007), thus inducing estrogenic effects such as hyperestrogenism in all animal species (Böhm and Razzai-Fazeli, 2005). In pigs, which are more sensitive than other species, ZEA is mainly biotransformed in the liver in α-Zearalenol (α-ZEA), which showed higher affinity to bind to estrogen receptors than ZEA and β-Zearalenol (β-ZEA) (Malekinejad et al., 2007). Common

clinical signs of ZEA intoxication are related to the hyperstimulation of estrogendependent tissues (Malekinejad et al., 2007). In gilts clinical signs include reddening and swelling of the vulva and an enlarged uterus, whereas in cycling sows fertility is impaired (Malekinejad et al., 2007; Minervini and Dell'Aquila, 2008).

In ruminants, ZEA is converted in the rumen mainly into  $\alpha$ -ZEA (Fig. 2.4) and, to a less extent, in  $\beta$ -ZEA (Fig. 2.5) (Abidin and Khatoon, 2012; Winkler et al., 2014) but the rate of absorption of the more polar  $\alpha$ -ZEA is poor.

In cow, infertility, reduced milk production and hyperestrogenism have been reported (Minervini and Dell'Aquila, 2008).

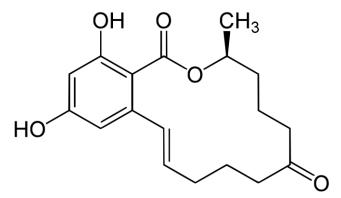


Figure 2.3: Chemical structure of zearalenone (ZEA).

**Figure 2.4:** Chemical structure of  $\alpha$ -zearalenol ( $\alpha$ -ZEA).

**Figure 2.5:** Chemical structure of  $\beta$ -zearalenol ( $\beta$ -ZEA).

## 2.2.6 Beauvericin

Beauvericin (BEA) (Fig. 2.6) is a mycotoxin firstly identified in the culture of the soil-borne entomopathogenic fungus Beauveria bassiana but also synthesized by several Fusarium spp. parasitic to important cereal grains such as corn, wheat, rice and barley (Leslie and Summerell, 2006). High contamination levels, up to 500 mg/kg of BEA, have been reported in commodities raising serious concerns about the potential impact of this fusariotoxin on human and animal health (Uhlig et al., 2007; Jestoi, 2008). BEA is a ionophoric molecule that can form stable and lipophilic complexes with cations and transport them into the lipophilic phase (Hilgenfeld and Saenger, 1982; Jestoi, 2008; Schoevers et al., 2016). According to this, the primary toxic action of BEA is considered to be related to its ionophoric properties. BEA is thus able to promote the transport of several cations through membranes disturbing the cell homeostasis (Hilgenfeld and Saenger, 1982; Jestoi, 2008; EFSA, 2014; Schoevers et al., 2016). It is well-established that an increase in the intracellular concentration of cations such as calcium and the subsequent activation of calcium-dependent endonucleases lead to DNA fragmentation which is related to several chronic diseases in humans and in animals with many different adverse health effects (Speijers and Speijers, 2004; Kouadio et al., 2007). In recent studies BEA was found to exert potent cytotoxicity against different cell lines (Jestoi, 2008; Ruiz et al., 2011; Prosperini et al., 2012; Mallebrera et al., 2016).

Figure 2.6: Chemical structure of beauvericin (BEA).

## 2.3 Fusarium mycotoxins and effects on reproductive function

#### 2.3.1 Fumonisins

The potential reproductive toxicity of fumonisins has been only recently investigated. Studies demonstrated that fumonisin B<sub>1</sub> (FB<sub>1</sub>) has the potential to impair fertility in pigs and rabbits delaying puberty and impairing semen quality and spermatogenesis (Ogunlade et al., 2006; Gbore and Egbunike, 2008; Ewuola and Egbunike, 2010). Moreover, *in vitro* studies showed that FB<sub>1</sub> affects some functional parameters of equine spermatozoa, such as sperm chromatin stability and motility (Minervini et al., 2010). Recently, FB<sub>1</sub> was found to directly affect *in vitro* porcine granulosa cell (GC) proliferation, steroid production and gene expression (Cortinovis et al., 2014). FB<sub>1</sub> at 10 µM decreased cell proliferation and strongly increased progesterone (P4) production thereby suggesting that this *Fusarium* mycotoxin may affect the normal follicle growth and oocyte survival and interfere with the endocrine regulation of the developing follicle in swine (Cortinovis et al., 2014).

Regarding a possible interaction between FB<sub>1</sub> and other Fusarium mycotoxins, a significant interaction between FB<sub>1</sub> at 10  $\mu$ M and  $\alpha$ -ZEA at 9.4  $\mu$ M was found on P4 production (Cortinovis et al. 2014). On the contrary, no significant interaction was

observed when GC were treated with FB<sub>1</sub> at 10  $\mu$ M combined with DON at 3.4  $\mu$ M (Cortinovis et al. 2014).

## 2.3.2 Deoxynivalenol

The potential of deoxynivalenol (DON) to act as endocrine disruptor has been investigated in previous studied and is the object of increasing interest (Medvedova et al., 2011; Han et al., 2016).

In previous studies, DON at 10  $\mu$ M was found to affect the process of follicular maturation with a decrease of the reserve pool of follicles resulting in a significant decrease in the number of normal follicles (Gerez et al., 2016). These results are in agreement with previous studies, where porcine cumulus-oocyte complexes exposed to DON at 0.02, 0.2, or 2  $\mu$ M showed an increase in cumulus cell death and degeneration, with a consequent significant reduction in the proportion of oocytes that reached metaphase II (Schoevers et al., 2010). Concerning the effect of DON on GC proliferation and steroidogenesis in pigs, Ranzenigo et al. (2008) reported DON increasing GC number at 0.034  $\mu$ M and 0.34  $\mu$ M and drastically reducing it at 3.4  $\mu$ M. Differently, Pizzo et al. (2016) found that DON did not alter cell proliferation of bovine GC at concentrations ranging from 0.1 to 3.3  $\mu$ M.

The effects of DON on steroidogenesis have been also investigated with the bovine GC model (Pizzo et al., 2015, 2016). DON, in presence of FSH and IGF1, was found to inhibit E2 release at concentrations ranging from 0.1 to 3.3 µM and P4 production at 0.33 and 3.3 µM (Pizzo et al., 2015). In absence of IGF1, DON at 3.3 µM significantly up-regulated CYP19A1 mRNA abundance, but had no effect on CYP11A1 mRNA abundance in bovine GC (Pizzo et al., 2016). These results seem to support the theory that DON promotes stability of several mRNAs interfering with post-transcriptional processes and avoiding their rapid degradation with several adverse effects on steroidogenesis in cattle (Pizzo et al., 2015; Pizzo et al., 2016).

#### 2.3.3 Zearalenone

It is already well established that zearalenone (ZEA) and its metabolites have strong estrogenic activities, being able to cause alteration in the reproductive tract (Fink-Gremmels, 1999; Malekinejad et al., 2007; Denli et al., 2017). Particularly, ZEA can induce estrogenic effects such as hyperestrogenism, anoestrus, ovarian atrophy and changes in the endometrium (Böhm and Razzai-Fazeli, 2005; Malekinejad et al., 2007; Minervini and Dell'Aquila, 2008). The effect of ZEA depends on several factors

including the reproductive status (prepuberal, cycling or pregnant) of the affected animal and the administration time and dose (Price et al., 1993; Tiemann and Dänicke, 2007; Döll and Dänicke, 2011; Holda and Glogowski, 2014). A correlation between the level of ZEA in mg/kg and the length of anestrus in days was found by Young and King (1986) who observed an increase of the weaning-to-estrus interval when increased ZEA was fed.

Döll et al. (2004) reported that after 5 weeks of feeding piglets in prepuberal stuatus with feed contaminated by ZEA up to 0.42 mg/kg, the mean weight of the uteri was significantly increased. ZEA and its metabolite adverse effects on estrus were also reported in a recent study (Daia et al., 2016). This *in vivo* study provided evidence that ZEA at 1.04 mg/kg accelerated the development of the ovaries in post-weaning piglets confirming that a diet contaminated by ZEA can accelerate the development of ovarian follicles in post-weaning piglets possibly leading to subsequent reproductive disorders (Daia et al., 2016). In another study, Minervini et al. (2006) studied the effects of ZEA and its metabolites with an *in vitro* culture system of equine GC. The results of this study showed GC apoptosis after exposure to ZEA at 0.1 μM. A recent study confirmed that ZEA induces necrosis and GC death in a dose-dependent manner via a caspase-3- and caspase-9-dependent mitochondrial pathway (Zhu et al. 2012).

The effects of α-ZEA and β-ZEA, ZEA metabolites, on pig oocytes have been investigated (Alm et al., 2002). Oocyte maturation rate resulted in a significant decrease when oocytes were exposed for 48 h to α-ZEA at concentrations up to 7.5 μM (Alm et al., 2002). Differently, β-ZEA showed a significant effect only at 30 μM (Alm et al., 2002). In a subsequent study, Tiemann et al. (2003) found α-ZEA and β-ZEA at concentrations of 15 and 30 µM to inhibit in vitro the FSH stimulated P4 synthesis in porcine GC (Tiemann et al., 2003). The ability of ZEA metabolites to affect steroid production in pig GC has also been reported by Ranzenigo et al. (2008) and Cortinovis et al. (2014). In both studies α-ZEA primarily increased progesterone (P4) production induced by FSH and IGF1, whereas estradiol (E2) production exhibited a biphasic dose-response to α-ZEA in the study conducted by Ranzenigo et al. (2008) and was not affected in the study of Cortinovis et al. (2014). Specifically, α-ZEA at 0.094 μM and 9.4 μM increased and decreased E2 production, respectively (Ranzenigo et al., 2008), whereas no effects on E2 production was reported after exposure to α-ZEA at 9.4 μM (Cortinovis et al., 2014). In comparison to monogastric, ruminants seem less susceptible to ZEA toxicity (Upadhaya et al., 2010; Pizzo et al., 2016). In rumen ZEA is converted into α-ZEA and β-ZEA, even if ZEA also undergoes hepatic biotransformation (Seeling et al., 2006). In cows the poor rate of absorption of the ZEA metabolites explains why clinical signs of hyperestrogenism are observed scarcely (Fink Gremmels, 2008). Estrogenic effects of ZEA in cows were found only after the ingestion of highly contaminated feed or after long-term exposure to contaminated feed materials (Fink Gremmels, 2008). In a recent study Pizzo et al. (2016) determined the impact of  $\alpha$ -ZEA and  $\beta$ -ZEA on GC function evaluating cell proliferation, steroid production and gene expression using a bovine GC model. Based on the results reported, in absence of IGF1,  $\alpha$ -ZEA at 3.1  $\mu$ M had inhibitory effects on cell proliferation, whereas it was found to inhibit both E2 and P4 production in GC at concentration ranging from 0.09 to 3.1  $\mu$ M in presence of IGF1 (Pizzo et al., 2016). Regarding  $\beta$ -ZEA, an inhibitory effect on cell numbers was found at 31  $\mu$ M both in presence and absence of IGF1, while E2 and P4 production was increased in the absence IGF1. The results obtained by Pizzo et al. (2016) on cell proliferation are in agreement with previous studies on pig GC conducted by Tiemann et al. (2003) and Ranzenigo et al. (2008) which demonstrated the adverse effects of  $\beta$ -ZEA. Regarding steroidogenesis previous studies on pigs (Ranzenigo et al., 2008) showed an increase in E2 production at 0.09  $\mu$ M whereas in cattle no results were obtained at the same concentration (Pizzo et al., 2016), suggesting a species-specific effect.

In relation to the possible interaction between ZEA metabolites and gene expression Pizzo et al. (2016) demonstrated that  $\alpha$ -ZEA in the presence of IGF1 did not affect CYP11A1 and CYP19A1 mRNA abundance. However, in previous studies on pigs Tienmann et al. (2003) showed that  $\alpha$ -ZEA at 5  $\mu$ M was able to increase CYP11A1 protein expression, whereas Ranzenigo et al. (2008) showed that  $\alpha$ -ZEA at 9.4  $\mu$ M decreased CYP11A1 mRNA abundance in porcine GC.

#### 2.3.4 Beauvericin

Beauvericin (BEA) is considered an emerging mycotoxin (Uhlig et al., 2007; Jestoi, 2008; EFSA, 2014). The ability of BEA to increase the cytoplasmic cation concentration may play an important role in the induction of cell apoptosis (Jow et al., 2004). Contaminations, up to 500 mg/kg of BEA, have been reported in commodities raising serious concerns about the potential impact of this mycotoxin (Jestoi, 2008). In previous studies BEA was found to exert potent cytotoxicity on pig, rodent and human cell lines (Klaric et al., 2006; Ruiz et al., 2011; Prosperini, et al., 2012). Recent studies, with estrogen, androgen, progestagen and glucocorticoid reporter gene assays (RGAs), demonstrated that BEA has the potential to modulate the endocrine system by antagonism of nuclear receptor transcriptional activity (Fernández-Blanco et al., 2016). Information on possible reproductive effects of BEA in domestic animals is lacking. In a recent study Schoevers et al. (2016) showed that BEA reduced the developmental competence of both the maturing oocyte and the two-four cell stage embryo in pigs, and that BEA only affected the rate of developing embryos. The authors exposed the cumulus-oocyte-complexes and developing embryos to BEA at concentrations ranging

from 0.31 to  $10~\mu\mathrm{M}$  and they studied the effects of this mycotoxins on viability, progesterone synthesis and apoptosis (Schoevers et al., 2016). As reported in this study BEA was toxic to embryos, oocytes and cumulus cells at concentrations exceeding  $0.5~\mu\mathrm{M}$ .

## 2.3.4.1 Granulosa cells in vitro model

The function of granulosa cells (GC) is essential in the process of folliculogenesis and oocyte growth and development (Petro et al., 2012). In fact GC are crucial in the delivery of nutrients to the oocyte and they play an important role in the ovarian steroidogenesis (Scaramuzzi et al., 2011). The use of the GC model to investigate the molecular mechanisms of several compounds, their adverse effects and the mechanisms underlying the process of ovarian follicular atresia has been already reported (Jolly et al., 1994; Kwintkiewicz et al., 2010). For its characteristics the GC model provides the opportunity to examine the factors that influence oocyte competence in a way not previously accessible (Dias et al., 2014). As reported by several authors the bovine GC model is not only able to reproduce the in vivo situation, but also allows comparable studies with humans (Anderiesz et al., 2000). Anderiesz et al. (2000) reported the response of bovine and human oocytes to pure recombinant preparations of human follicle stimulating hormone (FSH) and luteinizing hormone (LH) for meiotic maturation and subsequent developmental competence in vitro. Specifically, no significant difference was observed in terms of maturation of oocyte to metaphase II and embryonic development between bovines and humans (Anderiesz et al., 2000). These results are in agreement with subsequent studies where similarities in terms of embryonal genome activation and duration of preimplantation development were found, confirming that the bovine GC model represents a good in vitro model to evaluate the effects of contaminants in both bovines and humans (Petro et al., 2012).

## 2.4 Fusarium mycotoxins and in vitro intestinal models

#### 2.4.1 In vitro intestinal models

The intestinal tract represents the first interface between food and the internal body and is the primary target of dietary compounds, thus *in vitro* intestinal models are of great interest for several toxicological studies (Meca et al., 2011). The intestinal epithelial monolayer consists of several subsets of epithelial cells that constitute a physical and

biochemical network for the maintenance of the homeostasis in the gastrointestinal tract (Goto and Kiyono, 2012). The main functions of the intestinal epithelium include the protection of the body against potentially toxic compounds or microorganisms, and the prevention of the loss of important compounds such as water and solutes (Gordon et al., 2015). There are several *in vitro* intestinal models that can mimic oral toxicity with different advantages, limitations, issues and needs (Table 1).

Tabel 1: Models for intestinal absorption (Gordon et al., 2015)

	Models for intestinal absorption				
	Examples	Advantages	Limitations	Issues and needs	Мето
Cell-based systems	Caco-2 monolayer (e.g. CacoReady, Advancell)	- easily accessible - regulatory acceptance for BCS classification - reproducible - metabolically competent - information on active transport - Papp useful PBPK or bioavailability screening (e.g. BCS class)	- poor correlation to in vivo rat absorption data (since, e.g. no available human absorption data for pesticides)  - technical limitations (sufficient water solubility needed, dependence on shipment)	- definition of acceptance criteria (e.g. applicability domains, reference datasets, integrity tests,?)  - higher standardization of the method including standardized integrity tests (SOP?)  - think about new approaches (new in vitro models? combination of both?) that cover all pathways  - no reacceptance criteria in the regul conte data regul conte data regul content of them the regul content of them the regul content of them them them the data from the standardization of the standardized integrity tests (SOP?)  - think about new approaches (new in vitro models? combination of transport of the standardized in silico models? extent activity	
	T84 polarized cells	- depicts apical and basal surface - reflects villi like structures - well defined cellular junctions - easy to assess TEER for permeability measurements	- 7-10 days for polarization - technical limitations		- clarify transporter activity and extent of activity in Caco-2 model
Artificial surrogates	GiT-PAMPA (Pion)	- easily accessible - standardized - reproducible - Papp useful for PBPK	- only passive diffusion - no metabolism - poor correlation to <i>in vivo</i> rat absorption data - technical limitations (sufficient watersolubility, UV-activity needed)	mixtures	

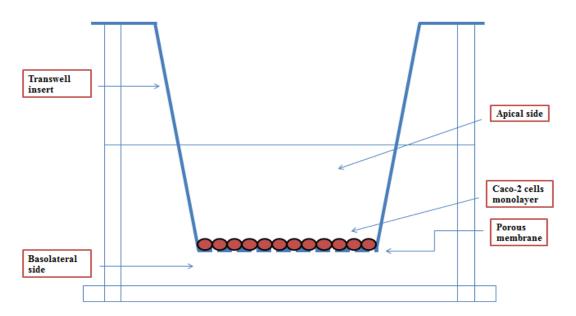
Among the intestinal models the most common are represented by immortalized human adenocarcinoma cell lines such as Caco-2 or T84 (Raffatellu et al., 2005; Khare et al., 2009; Tran et al., 2010). Caco-2 and T84 models are common to many studies, however each epithelial barrier model has specific characteristics for application (Ward and Tse, 1999). For example, it is well known that there are some significant differences between polarized Caco-2 and T84 cells. Caco-2 cell monolayers exhibit lower Trans-Epithelial Electrical Resistance (TEER) values on confluence than T84 cells (Gordon et al., 2015). An increase in the permeability of intestinal epithelial cells is correlate to a decrease in TEER values and this parameter has been used as an indicator of early sub-lethal epithelial toxicity (McCall et al., 2009; Gordon et al., 2015). Moreover, T84 cells differ from Caco-2 cells in Na<sup>+</sup>-dependent nucleoside transport systems (Ward and Tse, 1999). Other differences between these cell lines have been described before by several authors (McCool et al., 1990; Ward and Tse, 1999; Gordon et al., 2015). For example, compared to Caco-2 cells, T84 cells produce mucin in culture and are able to respond to external stimuli to exert innate immune responses thus mimicking more closely the intestinal surface conditions (McCool et al., 1990; Ward and Tse, 1999; Ou et al., 2009; Gordon et al., 2015). Culture-related conditions were shown to influence the expression of Caco-2 cell activity (Delie and Rubas, 1997; Sambuy et al., 2005; Turco et al., 2011). In order to reduce the variability, the TC7 clone was obtained from a late passage of the parental Caco-2 line (Turco et al., 2011). TC7 cells have demonstrated to consist of a more homogeneous population with more developed intercellular junctions, however Turco et al. (2011) reported the non-suitability of TC7 cells to predict intestinal absorption of highly lipophilic or poorly absorbed compounds (Turco et al., 2011).

Previous studies reported that IPEC-1 and IPEC-J2 cell lines are considered for their suitability good *in vitro* intestinal models (Lu et al., 2002; Schierack et al., 2006). Both cell lines are undifferentiated and derived from small intestine of piglets, however, while IPEC-1 derived from the small intestine, IPEC-J2 derived from mid-jejunum (Berschneider, 1989). IPEC-J2 cells grown in monolayers were first employed in transportant and cellular proliferation studies (Kandil et al., 1995; Rhoads et al., 1997). Neither of these cell lines is immortalized and therefore they are considered better models of normal porcine intestinal epithelium than transformed cell lines (Koh et al., 2007).

#### 2.4.2 Caco-2 in vitro intestinal model

Caco-2 cells, originally isolated from a human colorectal adenocarcinoma, represent a very well characterized *in vitro* model of epithelial barrier for intestinal absorption and metabolism studies (Fogh et al., 1977; Gilman and Cashman, 2006; Meca

et al., 2011; Prosperini et al., 2012). This cell line represents a good cell culture model for the study of absorption and metabolism in the small intestine, as well as the most used model for pharmacological and toxicological studies (Delie and Rubas, 1997; Le Ferrec et al., 2001; Sambuy et al., 2005). To better mimic the in vivo conditions of the intestine, Caco-2 cells are cultured on permeable filter supports which allow access of ions and nutrients to both sides of the cell monolayer (Turco et al., 2011). Caco-2 cells demonstrated to be highly dependent on culture conditions including medium composition and pH, seeding density and substrate nature (Ranaldi et al., 2003). However, this cell line is considered a suitable physiological model for studies of toxicity and transport of nutrients, cations and contaminants (Hidalgo et al., 1989; Artursson et al., 2001; Martel et al., 2001; Caloni et al., 2006; Caloni et al., 2012) (Fig. 2.7). When grown on a permeable filter support for 21 days Caco-2 cells are able to polarize and differentiate according to some typical enterocytic pathway, with apical microvilli and a basolateral surface, similar to the cellular surface in contact with intestinal vascular and lymphatic circulation (Pinto et al., 1983; Artursson et al., 2001; Caloni et al., 2012; Ferruzza et al., 2012). The Caco-2 cell full differentiation and polarization process has been associated with Trans-Epithelial Electrical Resistance (TEER) values exceeding 300  $\Omega$  cm<sup>2</sup> (Van Breemen and Li, 2005). In fact after confluence, TEER and permeability of marker molecules are usually used to investigate the integrity of the epithelial barrier tight junctions (Pinton et al., 2009). Specifically, TEER quantifies ion movement across cellular barriers and is considered a good indicator of the integrity of the epithelial barrier (De Angelis and Turco, 2011).



**Figure 2.7:** Transwell insert which separate the Apical (Ap) compartment from the basolateral (Bl) compartment.

Caco-2 cells also express some transporters and efflux proteins as well as metabolic enzymes that are normally expressed in the gut (Sun et al., 2002; Sambuy et al., 2005). *In vitro* methods based on chemical transport across Caco-2 monolayers are at present the most frequently and successfully exploited procedures to investigate intestinal permeability in humans (Yee, 1997; Yamascita et al., 2000; Van Breemen and Li., 2005; Nigsch et al., 2007), giving good correlation with the fraction absorbed in humans (Van Breemen and Li., 2005; Nigsch et al., 2007).

Passive diffusion is not the only absorption mechanism of xenobiotics, the active transport and efflux systems play also an important role (Yang, 2013). There are many intestinal transporters expressed on the small intestine and among these there are two main groups: the efflux (ABC family) and uptake (SLC family) transporters (Liu et al., 2013). Specifically, ABC transporters are composed by P-glycoprotein (P-gp), multidrug resistance 1 (MDR1), multidrug resistance-associated protein 2 (MRP2) and breast cancer resistance protein (BCRP) (Liu et al., 2013), whereas among SLC transporters we can found the organic cation transporters (OCTs), novel organic cation transporters (OCTNs) organic anion-transporting polypeptides (OATPs) and H+/peptide cotransporter (PEPT1) (Liu et al., 2013).

## 2.4.3 Fusarium mycotoxins and in vitro studies with intestinal barrier

Different *in vitro* models have been used to evaluate the effects of *Fusarium* mycotoxins on the intestinal barrier (Table 2). Table 2 summarizes the cell models used to assess the impact of the principal Fusariotoxins on the intestinal epithelium.

Tabel 2: Fusarium mycotoxins and in vitro studies with intestinal models

Mycotoxin	Model	References
	Caco-2 cells	Stevens et al., 1997; Caloni et al., 2002; De Angelis et al., 2005; Kouadio et al., 2005; Kouadio et al., 2007; Fernández-Blanco et al., 2016; Romero et al., 2016.
$FB_1$	Ipec-1	Bouhet et al., 2004; Bouhet and Oswald, 2007; Loiseau et al., 2007
	Ipec-J2	Goossens et al., 2012; Wang et al., 2013
	HT-29	Minervini et al., 2014
	SW742	Mahamoodi et al., 2012
BEA	Caco-2 cells	Fernández-Blanco et al., 2016
DEA	Ipec-1	Springler et al., 2016
	Caco-2 cells	Kasuga et al., 1998; Kouadio et al., 2005; Sergent et al., 2005; Manda et al., 2015; Vejdovszkya et al., 2016;
DON	Ipec-1	Pinton et al., 2009; Pinton et al., 2010; Diesing et al., 2011; Alassane-Kpembi et al., 2015
	Ipec-J2	Gu et al 2014; Broekaert et al, 2016; Gu et al 2016
	Caco-2 cells	Kouadio et al 2005; Kouadio et al, 2007; Videmann et al 2008; Gao et al., 2016
ZEA	Ipec-1	Wan et al, 2013b; Taranu et al, 2014
	Ipec-J2	Goossens et al, 2012; Wan et al., 2013b

#### 2.5 References

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3. In vitro effects of fumonisin  $B_1$  alone and combined with deoxynivalenol,  $\alpha$ -zearalenone,  $\beta$ -zearalenone and beauvericin on bovine granulosa

### 3.1 Introduction

Mycotoxins, secondary metabolites produced by moulds, frequently occur in food and feed (Oswald et al 2005; Schoelleberger et al., 2007) and a worldwide contamination is related to *Fusarium* mycotoxins (Scott, 1997; Placinta et al., 1999; Schollenberger et al., 2007; Lombaert et al., 2003).

Fusarium mycotoxins can be classified in two groups: i) the "traditional" mycotoxins including fumonisins, trichothecenes and zearalenone (ZEA) and ii) the so-called "emerging" mycotoxins such as beauvericin (BEA), fusaproliferin, enniatins (ENNs) and moniliformin (Jestoi, 2008).

Among the fumonisins, fumonisin B<sub>1</sub> (FB<sub>1</sub>) is considered the main toxin in terms of toxicity (Rodrigues and Naehrer, 2012) and its co-occurrence with other *Fusarium* mycotoxins is well established (EFSA, 2005; Voss et al., 2007). Leukoencephalomalacia in horses (Kellerman et al., 1990; Raymond et al., 2005), pulmonary edema in pigs (Harrison et al., 1990; Tiemann and Danick, 2007) and nephropathy in rabbits (Gumprecht et al., 1995; Wangikara et al., 2005) are species-specific effects related to the exposure to FB<sub>1</sub>. Cattle seem less sensitive than other animals such as horses or pigs to FB<sub>1</sub> (Osweiler et al., 1993; Mathur et al., 2001; Tiemann and Danick, 2007; Fink-Gremmels, 2008).

The complex mechanism of action of FB<sub>1</sub> (Wang et al., 1991; Voss et al., 2002; Voss et al., 2007; Gbore et al., 2012) is based on the inhibition of ceramide synthase (Wang et al., 1991; Voss et al., 2007; Luongo et al. 2008) with the alteration of the sphingolipid metabolism (Wang et al., 1991; Wang et al; 1992; Merrill et al., 2001; Marasas et al., 2004; Voss et al., 2007; Smith, 2012).

In relation to reproductive effects, no data are available on cattle, while *in vitro* studies demonstrated that FB<sub>1</sub> also combined with DON or  $\alpha$ -ZEA, affects porcine granulosa cells (GC) (Cortinovis et al., 2014).

Among trichothecenes, DON, also called vomitoxin for its emetic effects, is one of the most important and widespread in cereal grains and animal feed (Rotter et al., 1996). The co-occurrence of DON with other mycotoxins such as ZEA is a common feature of several studies (Richard, 2007; Pinton and Oswald, 2014).

Feed refusal was observed in pigs and in cattle (Trenholm et al., 1984; Dersjant-Li et al., 2003; Akande et al., 2006; Rodrigues and Naehrer, 2012) after exposure to DON.

Reproductive disorders have also been reported in livestock (Diekman and Green, 1992), but little is known about DON effects on ruminant reproduction (Seeling et al., 2006).

ZEA is a non-steroidal estrogenic Fusarium mycotoxin known to contaminate various crops, especially corn, wheat, barley and oats (Cheeke, 1998; Zinedine et al., 2007;

Ferrigo et al., 2016). ZEA is able to bind to cytosolic estrogen receptors in target cells (Riley and Norred, 1996) and acts like endocrine disruptor with estrogenic effects (Frizzell et al., 2011). Its activity is related to the transcription of estrogen-responsive genes and consequently to the modulation of the translation of new proteins and the expression of estrogenic effects upon the target cells (Parveen et al., 2009).

The hyperstimulation of estrogen-dependent tissues is the main effect of ZEA exposure (Böhm and Razzai-Fazeli, 2005; Malekinejad et al., 2007). The  $\alpha$ -ZEA metabolite seems to have an higher binding affinity to estrogen receptors compared to ZEA and  $\beta$ -ZEA metabolite (Fink-Gremmels, 2008).

As reported for FB<sub>1</sub>, cattle are considered to be less susceptible than other domestic animals to the adverse effects of ZEA (Upadhaya et al., 2010). In rumen ZEA is converted into α-ZEA and β-ZEA (Abidin and Khatoon, 2012; Winkler et al., 2014) and the rate of absorption of the more polar α-ZEA is poor and hyprestrogenism is infrequently observed in cows. Study conducted on liver microsomes demonstrated that the ratio between α-ZEA and β-ZEA formation varies among animal species and that in ruminants, because ZEA undergoes hepatic metabolism, β-ZEA seems to prevail as results of hepatic metabolism, whereas α-ZEA is the most frequent metabolite in pigs (Malekinejad et al., 2007; Fink-Gremmels, 2008). ZEA and its metabolites exert their toxicity by binding to estrogens receptors and the clinical signs of hyperestrogenism in ruminants are generally observed after long-term exposure to ZEA and after ingestion of highly contaminated feed (Fink-Gremmels, 2008; Upadhaya et al., 2010). ZEA exposure has also been linked to infertility, reduction in milk production, inhibition of oocytes maturation to metaphase II and an alteration of steroidogenesis in cattle (Minervini and Dell'Aquila, 2008; Pizzo et al., 2016).

BEA is a cyclic hexadepsipeptide that was first isolated from the culture of the soil-borne entomopathogenic fungus *Beauveria bassiana* (Leslie and Summerell, 2006). BEA is also synthesized by several *Fusarium* spp. parasitic to important cereal grains including corn, wheat, rice and barley (Leslie and Summerell, 2006). High contamination levels, up to 500 mg/kg of BEA, detected in surveys carried out across Europe are raising serious concerns about the potential impact of BEA on animal and human health (Uhlig et al., 2007; Jestoi, 2008). Few data are available on BEA toxicity so far (EFSA, 2014). BEA proved to exert potent cytotoxicity against several mammalian cell lines (Jestoi, 2008; Klarić et al., 2008; Ferrer et al, 2009; Ruiz et al., 2011; Prosperini et al., 2012; Mallebrera et al., 2016) and this effect seems to be related to its ionophoric properties (EFSA, 2014). Accordingly, BEA is able to promote the transport of cations, such as calcium, through the membranes disturbing their normal physiological concentrations in the cell and thus affecting ionic homeostasis (Jestoi, 2008).

To date information on the effects of FB<sub>1</sub> alone or combined with other Fusarium mycotoxins on reproduction in cattle is lacking.

In the present study primary bovine GC, which are considered a reliable *in vitro* model for reproductive toxicological research (Petro et al., 2012), were used to evaluate the individual and combined effects of FB<sub>1</sub> with DON,  $\alpha$ -ZEA,  $\beta$ -ZEA and BEA on cell proliferation, steroid production and gene expression.

#### 3.2 Materials and methods

#### 3.2.1 Reagents and hormones

Reagents were: Dulbecco's Modified Eagle Medium (DMEM), Ham's F12, Fumonisin B<sub>1</sub> (FB<sub>1</sub>), Deoxynivalenol (DON), α-Zearalenol (α-ZEA) β-Zearalenol (β-ZEA) and Beauvericin (BEA) obtained from Sigma Chemical Co. (St. Louis, MO); fetal calf serum (FCS) obtained from Atlanta Biologicals (Flowery Branch, GA); purified ovine follicle stimulating hormone (FSH; FSH activity, 15 NIH-FSH-S1 U/mg) obtained from Dr. A. F. Parlow, National Hormone and Pituitary Program (Torrance, CA); recombinant human insulin-like growth factor 1 (IGF1) obtained from R&D Systems (Minneapolis, MN); and testosterone obtained from Steraloids (Wilton, NH).

#### 3.2.2.Cell culture

Ovaries from non-pregnant beef cows were collected from a slaughterhouse as previously described (Langhout et al., 1991; Lagaly et al., 2008). Based on surface diameter small follicles (1-5 mm) were aspirated (Fig. 3.1) and GC were recovered from follicular fluid by centrifugation (291 x g for 10 min). GC were washed three times with 7 mL of serum-free medium and resuspended in 2 mL of enzyme containing medium (0.5 mg/mL of DNase and 1.25 mg/mL of collagenase) to prevent clumping of cells as previously described (Spicer et al., 2002; Lagaly et al., 2008). Numbers of viable cells were determined using the trypan blue exclusion method (Langhout et al., 1991; Spicer et al., 1993; Tiemann et al., 2003). Viable cells (2.5 x 10<sup>5</sup> in 20-80 µL of medium) were plated in 24-well Falcon multiwell plates (Becton Dickinson, Lincoln Park, NJ, USA) in 1 mL of basal medium composed of a mixture of 1:1 DMEM and Ham's F-12 containing glutamine (2 mM), gentamicin (0.12 mM) and sodium bicarbonate (38.5 mM). Cultures were kept at 38.5 °C in a humidified 95% air and 5% CO2 environment and medium was changed every 24 h. To obtain an optimal attachment, cells were maintained in the presence of 10% FCS for the first 48 h of culture. After this time, GC were washed twice with serum-free medium (0.5 mL) and the various treatments applied

in serum-free medium containing 500 ng/mL of testosterone (as an estradiol precursor) for 48 h with a medium change after 24 h.



Figure 3.1: Collection of granulosa cells from small bovine follicles (1-5 mm) via needle aspiration.

## 3.2.3 Determination of GC numbers

Medium was collected from individual wells and frozen at -20 °C for subsequent steroid analyses. Numbers of GC, in the same wells from which medium was collected, were determined by a Coulter counter (model Z2; Beckman Coulter, Inc., Hialeah, FL) (Fig. 3.2) as previously described (Lagaly et al., 2008), and used to calculate steroid production on ng or pg per 10<sup>5</sup> cell basis. Briefly, cells were gently washed twice with 0.9% saline solution (500 mL), exposed to 500 mL of trypsin (0.25% wt/vol; 2.5 mg/mL) for 20 min at room temperature, and then scraped from each well and enumerated as previously described (Langhout et al., 1991; Lagaly et al., 2008).



Figure 3.2: Determination of granulosa cell numbers using a Z2 Coulter® Particle Count and Size Analyzer.

## 3.2.4 Determination of steroid concentrations

Concentrations of progesterone (P4) and estradiol (E2) in culture medium were determined by radioimmunoassay (RIA) as previously described (Spicer and Chamberlain, 1998; Lagaly et al., 2008). The intra- and inter-assay coefficients of variation were 7% and 13%, respectively for the P4 RIA, and 8% and 17%, respectively for the E2 RIA. P4 production was evaluated because it, like E2 production, increases as GC undergo differentiation (Hsueh et al., 1984; Ainsworth et al., 1990).

#### 3.2.5 Progesterone RIA

Progesterone RIA were conducted using rabbit antiserum (X-16), which serves as the first antibody (diluted 1:3000 with assay buffer: PBS, EDTA, NaN3, and gelatin), raised against BSA-11 glutamate derivative as described by Baraño and Hammond (1985). Goat anti-rabbit antibody (diluted 1:15 with assay buffer) was used as the second antibody (Linco Research, Inc., St. Charles, MO). [125I]Iodo-progesterone (ICN Biomedicals, Costa Mesa, CA) was used as the tracer. A progesterone standard curve was prepared from a stock concentration of 80.0 ng/mL that was serially diluted with assay buffer to concentrations of 40.0, 20.0, 10.0, 5.0, 2.5, 1.25, 0.625, 0.31, and 0.16 ng/mL. In duplicate, 20 μL to 100 μL of medium samples were combined with the appropriate volume of assay buffer to make a total volume of 100 μL. One hundred μL

of tracer and first antibody were added and all samples were mixed and allowed to incubate at 37 °C for 1 h. Following incubation, 200 µL of second antibody were added and all samples were incubated overnight at 4 °C. The following day, 50 µL of normal rabbit serum (NRS) (diluted 1:5 with assay buffer from a 15% NRS stock) were added to all samples. Samples were centrifuged at 4 °C in a Sorvall Model RC-3 (Thermo Fisher Scientific, Inc., Miami, OK) at 1800 x g for 25 min. Supernatant was aspirated and precipitates were counted for 1 min using a Cobra AII Auto-Gamma counter (Packard Instrument Co., Downers Grove, IL). The intra- and interassay coefficients of variation were 7 and 13%, respectively, for the progesterone RIA.

#### 3.2.6 Estradiol RIA

Estradiol RIA were conducted using anti-estradiol rabbit antibody (diluted 1:12 with assay buffer), which serves as the first antibody (Lilly Research Laboratories, Indianapolis, IN), and goat anti-rabbit antibody (diluted 1:15 with assay buffer) which serves as the second antibody (Linco Research, Inc., St. Charles, MO). Radiolabeled estradiol (125I-estradiol) was used as the tracer (ICN Biomedicals, Costa Mesa, CA). The assay buffer was the same as the progesterone RIA buffer described above. An estradiol dose response curve was prepared from a stock concentration of 256 pg/100 µL that was serially diluted to 128, 64, 32, 16, 8, 4, 2, 1, and 0.5 pg/100  $\mu$ L using assay buffer. In duplicate, sample media was added at either 50 or 100 µL and (if needed) combined with assay buffer to make a total volume of 100 µL. Two hundred µL of tracer were added to all samples, along with 100 µL of first antibody. All tubes were mixed and allowed to incubate for 1 h at 37 °C. Following this incubation, 200 µL of second antibody were added and the assay allowed to incubate at 4 °C overnight. The following day, assay tubes were centrifuged, supernatant aspirated, and precipitate counted as described for the progesterone RIA. The intra- and interassay coefficients of variation were 8% and 17%, respectively, for the estradiol RIA.

#### 3.2.7 RNA extraction

At the end of the treatment period, cells from two replicate wells were lysed in 500 µL of TRIzol® reagent and RNA was extracted as previously described (Voge et al., 2004; Aad et al., 2006). Briefly, 250 µL TRIzol® reagent was added to all wells and cells were lysed by repeated pipetting and then combined with their respective replicates. Combined wells were then transferred to 1.5 mL eppendorf tubes. Each treatment containing 4 wells generated 2 replicate samples of RNA. Cell lysates were incubated in

TRIzol® reagent for approximately 5 min at room temperature, then 100 µL of chloroform was added to each sample followed by a 15 s vortex. After approximately a 2 min incubation at room temperature, samples were centrifuged at 3500 x g for 30 min at 4 °C using eppendorf centrifuge 5417C (Brinkmann Instruments, Westbury, NY). The upper aqueous phase of each sample was then transferred to a fresh eppendorf tube and RNA was precipitated using 250 µL isopropanol. Samples were incubated at room temperature for 10 min and then centrifuged at 3500 x g for 10 min at 4 °C. The RNA pellets were washed after discarding the supernatant with 500 µL of 70% ethanol and allowed to dry at room temperature. The RNA pellets were suspended in 16.5  $\mu L$  of DEPC-treated water. RNA was quantitated by spectrophotmetry at 260 nm using a NanoDrop ND-1000 spectrophotometer (NanoDrop Technologies, Wilmington, DE). Aliquots of 1.5 μL of RNA were used to determine the concentration in ng/μL as well as the purity given as a ratio of 260/280 nm where values between 1.8 and 2.2 were acceptable. RNA was then diluted to 10 ng/µL in DEPC-treated water and stored at -80 °C until used for quantification of target gene expression. Just prior to use, an RNA aliquot was thawed on ice for 3-5 min.

#### 3.2.8 Real-time PCR

The target gene primers (forward, reverse) and probe sequences for bovine enzyme (CYP19A1; Accession NM\_174305) aromatase were TGCCAAGAATGTTCCTTACAGGTA, CAGAGTGACCTTCATCATGACCAT and CATTTGGCTTTGGGCCCCGG, respectively; and for bovine P450 side-chain (CYP11A1; cleavage enzyme Accession NM\_176644) were CTCCGTGACCCTGCAGAGATAC, ATAGACGGCCACTTGTACCAATG and TTGGTTCTTCGAGATTACATGATTCCTGCC, respectively (Lagaly et al., 2008; Spicer and Aad, 2007). The differential expression of target gene mRNA in GC was quantified using the one-step multiplex real-time RT-PCR reaction for Tagman® Gold RT-PCR Kit (Applied Biosystems, Foster City, CA) as previously described (Spicer and Aad, 2007). All samples were run in duplicate. The 18S ribosomal RNA values were used as internal controls to normalize samples for any variation in amounts of RNA loaded, and relative quantification of target gene mRNAs was expressed using the comparative threshold cycle method as previously described (Voge et al., 2004; Aad et al., 2006). Briefly, the  $\Delta$ Ct was determined by subtracting the 18S Ct value from the target unknown value. For each target gene, the  $\Delta\Delta$ Ct was determined by subtracting the higher  $\Delta Ct$  (the least expressed unknown) from all other  $\Delta Ct$  values. Fold changes in target gene mRNA abundance were calculated as being equal to  $2^{-\Delta\Delta Ct}$ .

### 3.2.9 Experimental design

Experiment 1 was performed to evaluate the effects of FB<sub>1</sub> alone and combined with β-ZEA on GC proliferation and steroidogenesis. Cells were cultured for 48 h in 10% FCS, washed twice with serum-free medium as described earlier, and then treated for 48 h in serum-free medium containing testosterone (500 ng/mL), FSH (30 ng/mL) and IGF1 (30 ng/mL) with or without FB<sub>1</sub> at 0, 30, 100 ng/mL (0; 0.042; 0.13 μM) and β-ZEA at 0 or 30 ng/mL (0; 0.094 μM). After 48 h of treatment, cells were counted and medium was collected for E2 and P4 RIA. Doses of FSH and IGF1 were selected based on previous studies (Spicer et al., 2002; Ranzenigo et al., 2008). Because IGF1 alone has little or no effect on steroid production, FSH was added to all treatments (Spicer et al., 1993; 2002; Ranzenigo et al., 2008).

Experiment 2 was designed to evaluate the effects of interaction between FB<sub>1</sub>, DON and DON with  $\beta$ -ZEA. GC were cultured for 48 h in 10% FCS washed twice with serum-free medium as described earlier, and cells treated for 48 h in serum-free medium containing 500 ng/mL of testosterone, 30 ng/mL of FSH and 30 ng/mL of IGF1with FB<sub>1</sub> at 0, 30, 100 ng/mL (0; 0.042; 0.13  $\mu$ M), DON at 0 or 100 ng/mL (0; 0.33  $\mu$ M) and  $\beta$ -ZEA at 0 or 30 ng/mL (0; 0.094  $\mu$ M). After 48 h of treatment, medium was collected for P4 and E2 RIA, and cells were counted.

Experiment 3 was performed to evaluate the effects of FB<sub>1</sub> alone and combined with  $\alpha$ -ZEA and  $\beta$ -ZEA in the presence of FSH and with or without IGF1 on GC proliferation and steroidogenesis. Cells were cultured for 48 h in 10% FCS, washed twice with serum-free medium as described earlier, and cells treated for 48 h in serum-free medium containing testosterone (500 ng/mL), FSH (30 ng/mL) with or without IGF1 (30 ng/mL). FB<sub>1</sub> was tested at 5  $\mu$ g/mL (6.9  $\mu$ M) alone and combined with  $\alpha$ -ZEA at 5  $\mu$ g/mL (15.6  $\mu$ M) and  $\beta$ -ZEA at the same concentration (15.6  $\mu$ M). After 48 h cells were counted and medium was collected for P4 and E2 RIA.

Experiment 4 was carried out to compare the dose response of FB<sub>1</sub> and BEA on FSH plus IGF1–induced GC proliferation and steroidogenesis. GC were cultured for 48 h in 10% FCS washed twice with serum-free medium as described earlier, and cells treated for 48 h in serum-free medium containing testosterone (500 ng/mL), FSH (30 ng/mL) and IGF1 (30 ng/mL) with the various doses of FB<sub>1</sub> (i.e., 0, 0.3, 1, 3, 10 μM) or BEA (0, 0.3, 1, 3, 10 μM). After 48 h of treatment, medium was collected for determination of progesterone and estradiol concentrations via RIA, and cells were counted. Doses of FSH and IGF1 were selected based on previous studies (Spicer et al., 2002; Ranzenigo et al., 2008). Because IGF1 alone has little or no effect on steroid production, FSH was added to all treatments (Spicer et al., 1993; 2002; Ranzenigo et al., 2008).

Experiment 5 was performed to evaluate the effects of FB<sub>1</sub> and BEA, alone and in combination, on GC proliferation and steroidogenesis. Cells were cultured for 48 h in

10% FCS, washed twice with serum-free medium as described earlier, and cells treated for 48 h in serum-free medium containing testosterone (500 ng/mL), FSH (30 ng/mL) and IGF1 (30 ng/mL) with or without FB<sub>1</sub> (3 μM) and BEA (3 μM). After 48 h of treatment, cells were counted and medium was collected for determination of progesterone and estradiol concentrations via RIA.

Experiment 6 was designed to evaluate the dose response of FB<sub>1</sub> alone and combined with BEA on GC proliferation and steroidogenesis. Cells were cultured for 48 h in 10% FCS, washed twice with serum-free medium, and then treated for 48 h in serum-free medium containing testosterone (500 ng/mL), FSH (30 ng/mL) and IGF1 (30 ng/mL) with the various doses of FB<sub>1</sub> (i.e., 0, 0.5, 1, 1.5, 3, or 6 μM) with or without BEA (3 μM). After 48 h of treatment, medium was collected for determination of progesterone and estradiol concentrations via RIA, and cells were counted.

Experiment 7 was designed to determine the dose response of BEA alone and combined with FB<sub>1</sub> on GC proliferation and steroidogenesis. Cells were cultured for 48 h in 10% FCS, washed twice with serum-free medium, and then treated for 48 h in serum-free medium containing FSH (30 ng/mL) and IGF1 (30 ng/mL) with the various doses of BEA (i.e., 0, 0.5, 1, 1.5, 3, or 6  $\mu$ M) with or without FB<sub>1</sub> (3  $\mu$ M). After 48 h of treatment, medium was collected for determination of progesterone and estradiol concentrations via RIA, and cells were counted.

Experiment 8 was performed to determine the effects of FB<sub>1</sub> and BEA on *CYP11A1* and *CYP19A1* mRNA abundance. Cells were cultured for 48 h in 10% FCS, washed twice with serum-free medium as described earlier, and cells treated for 24 h in serum-free medium containing testosterone (500 ng/mL), FSH (30 ng/mL), IGF1 (0 or 30 ng/mL) and FB<sub>1</sub> (30  $\mu$ M) or BEA (30  $\mu$ M). After 24 h of treatment, medium was aspirated and cells were lysed for RNA extraction.

Experiment 9 was carried out to determine the effect of  $FB_1$  and BEA on serum-stimulated GC proliferation. Cells were cultured for 4 days in 10% FCS. During the last 2 days of culture, cells were treated as follows: control (no additions),  $FB_1$  (10  $\mu$ M) or BEA (10  $\mu$ M). At the end of treatment, cells were counted.

## 3.2.10 Statistical analysis

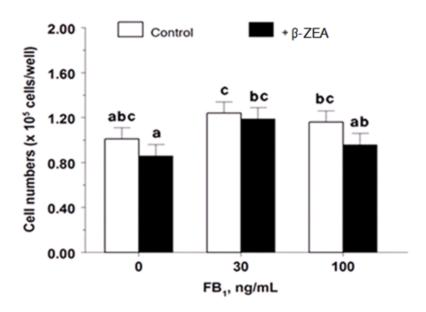
Experimental data are presented as the least squares means ± SEM of measurements from replicated experiments (n=3) with each treatment applied in triplicate or duplicate wells for each replicate experiment. Each replicated experiment was derived from a different pool of GC generated from a total volume of 10 mL follicular fluid obtained from twenty to thirty ovaries. For mRNA experiments, treatments were applied in quadruplicate culture wells with each mRNA sample being

obtained from two wells. Treatment effects were assessed by factorial ANOVA designs in the GLM procedure of the Statistical Analysis System (SAS) using SAS for Windows (version 9.2, SAS Institute Inc., Cary, NC). Steroid production was expressed as ng or pg/10<sup>5</sup> cells per 24 h, and GC numbers determined at the end of the experiment were used for this calculation. A P-value of less than 0.05 was considered statistically significant. Mean differences in steroid production, cell numbers and mRNA abundance between treatments were determined using the Fisher's protected least significant difference (LSD) procedure (Ott, 1977).

## 3.3 Results

3.3.1 Experiment 1: Dose response of FB<sub>1</sub> alone and combined with  $\beta$ -ZEA on GC numbers and steroid production in the presence of FSH with IGF1

The results revealed that FB<sub>1</sub> alone at 30 and 100 ng/mL (0; 0.042; 0.13  $\mu$ M) had no effect (P>0.10) on GC numbers. In the presence of  $\beta$ -ZEA at 30 ng/mL (0.094  $\mu$ M), FB<sub>1</sub> at 30 ng/mL (0.042  $\mu$ M) showed a stimulatory effect (P<0.05) on GC numbers (Fig. 3.3), whereas FB<sub>1</sub> at 100 ng/mL (0.013  $\mu$ M) increased (P<0.05) P4 production (Fig. 3.4). FB<sub>1</sub> at 30 ng/mL (0.042  $\mu$ M) and 100 ng/mL (0.13  $\mu$ M) was found to amplify the inhibitory effect of  $\beta$ -ZEA at 30 ng/mL (0.094  $\mu$ M) on E2 production (Fig. 3.5).



**Figure 3.3:** Effect of FB<sub>1</sub> on numbers of GC from bovine follicles. Means ( $\pm$  SEM) without a common letter (a–c) differ (P < 0.05). FB<sub>1</sub>, fumonisin B<sub>1</sub>;  $\beta$ -ZEA,  $\beta$ -zearalenone; SEM, standard error of the mean.

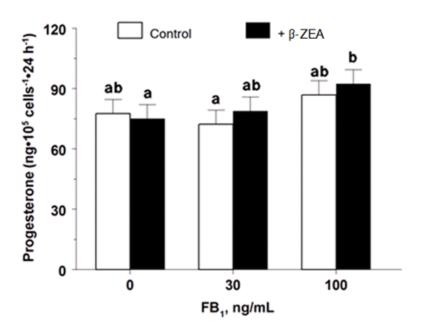


Figure 3.4: Effect of FB<sub>1</sub> on progesterone production of GC from bovine follicles. Means ( $\pm$  SEM) without a common letter (a–b) differ (P < 0.05). FB<sub>1</sub>, fumonisin B<sub>1</sub>;  $\beta$ -ZEA,  $\beta$ -zearalenone; SEM, standard error of the mean.

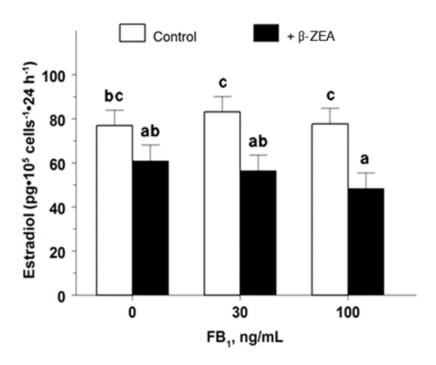


Figure 3.5: Effect of FB<sub>1</sub> on estradiol production of GC from bovine follicles. Means ( $\pm$  SEM) without a common letter (a–c) differ (P < 0.05). FB<sub>1</sub>, fumonisin B<sub>1</sub>;  $\beta$ -ZEA ,  $\beta$ -zearalenone;, SEM, standard error of the mean.

## 3.3.2 Experiment 2: Dose response of FB<sub>1</sub> alone and combined with DON and $\beta$ -ZEA on GC numbers and steroid production in the presence of FSH with IGF1

No significant interaction (P>0.10) existed between FB<sub>1</sub> at 30 ng/mL and 100 ng/mL (0; 0.042; 0.13  $\mu$ M) and DON at 100 ng/mL (0.33  $\mu$ M) or between FB<sub>1</sub> at 30 ng/mL and 100 ng/mL (0; 0.042; 0.13  $\mu$ M) and the combination of  $\beta$ -ZEA at 30 ng/mL (0.094  $\mu$ M) and DON at100 ng/mL (0.33  $\mu$ M) on cell numbers (Fig. 3.6). Regarding P4 and E2 production, no significant interaction (P>0.10) was observed between FB<sub>1</sub> at 30 ng/mL and 100 ng/mL (0; 0.042; 0.13  $\mu$ M) and DON at 100 ng/mL (0.33  $\mu$ M) alone or with  $\beta$ -ZEA at 30 ng/mL (0.094  $\mu$ M) on P4 (Fig. 3.7) or E2 (Fig. 3.8) production, but  $\beta$ -ZEA inhibited (P<0.05) P4 production in DON treated GC (Fig. 3.7).

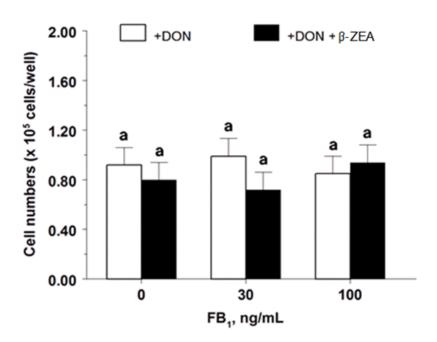


Figure 3.6: Interaction between FB<sub>1</sub> and DON or β-ZEA on proliferation of bovine GC. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a) differ (P < 0.05). FB<sub>1</sub>, fumonisin B<sub>1</sub>; DON, deoxynivalenol; β-ZEA, β-zearalenone; SEM, standard error of the mean.

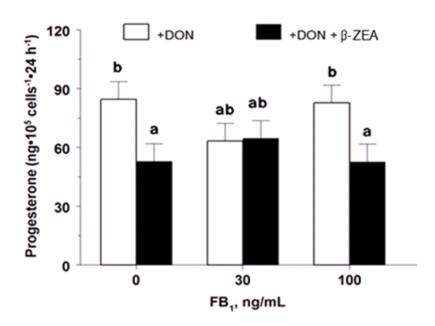


Figure 3.7: Interaction between FB<sub>1</sub> and DON or  $\beta$ -ZEA on progesterone production by GC from bovine follicles. Values are means  $\pm$ SEM from three separate experiments. Means without a common letter (a–b) differ (P < 0.05). DON, deoxynivalenol; FB<sub>1</sub>, fumonisin B<sub>1</sub>;  $\beta$ -ZEA,  $\beta$ -zearalenone; SEM, standard error of the mean.

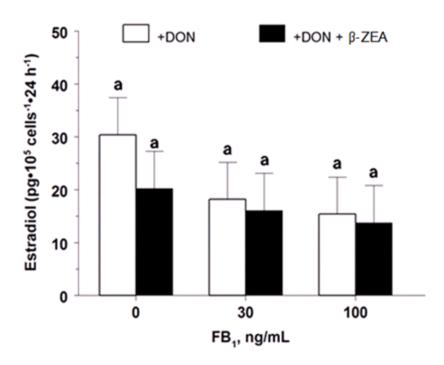
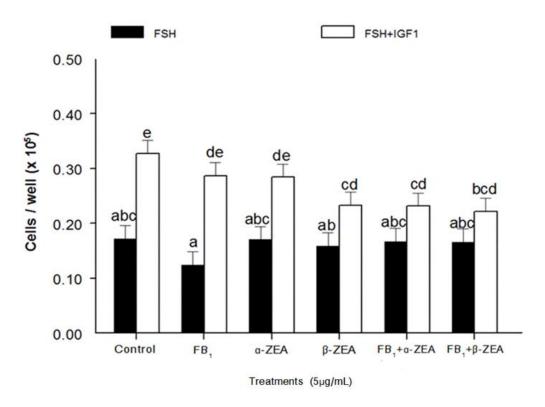


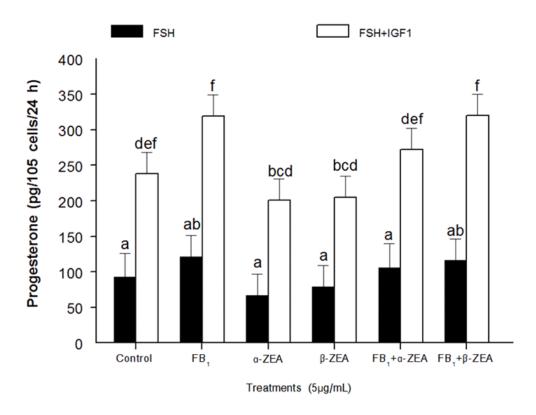
Figure 3.8: interaction between FB<sub>1</sub> and DON or  $\beta$ -ZEA on estradiol production by GC from bovine follicles. Values are means  $\pm$ SEM from three separate experiments. Means without a common letter (a) differ (P < 0.05). DON, deoxynivalenol; FB<sub>1</sub>, fumonisin B<sub>1</sub>;  $\beta$ -ZEA,  $\beta$ -zearalenone; SEM, standard error of the mean.

## 3.3.3 Experiment 3: Effect of FB<sub>1</sub> alone and combined with either a-ZEA or $\beta$ -ZEA on GC numbers and steroid production

Experiment 3 was conducted to assess the effects of FB<sub>1</sub> alone and combined with  $\alpha$ -ZEA and  $\beta$ -ZEA on cell proliferation and steroidogenesis with and without IGF1. In the absence of IGF1, GC numbers (Fig. 3.9), P4 production (Fig. 3.10), and E2 production (Fig. 3.11) were not affected by any of the mycotoxins or their combinations. IGF1 induced (P<0.05) GC proliferation and this increase was blocked (P < 0.05) after exposure to  $\beta$ -ZEA alone and FB<sub>1</sub> with either  $\alpha$ -ZEA or  $\beta$ -ZEA (Fig. 3.9). In contrast, IGF1-induced P4 production was not affected (P>0.10) by FB<sub>1</sub>,  $\alpha$ -ZEA ,  $\beta$ -ZEA or their combinations (Fig. 3.10). However, IGF1-induced E2 production was increased (P<0.05) with FB<sub>1</sub> alone (Fig. 3.11). In contrast,  $\alpha$ -ZEA alone,  $\beta$ -ZEA alone, or their combination with FB<sub>1</sub> decreased (P<0.05) IGF1-induced E2 production (Fig. 3.11).



**Figure 3.9:** Effect of FB<sub>1</sub> on numbers of GC from bovine follicles. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–e) differ (P < 0.05). FB<sub>1</sub>, fumonisin B<sub>1</sub>;  $\alpha$ -ZEA,  $\alpha$ -zearalenone;  $\beta$ -ZEA,  $\beta$ -zearalenone; SEM, standard error of the mean.



**Figure 3.10:** Interaction between FB<sub>1</sub>, α-ZEA or β-ZEA on progesterone production of bovine GC. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a-f) differ (P < 0.05). FB<sub>1</sub>, fumonisin B<sub>1</sub>; α-ZEA, α-Zearalenone;, β-ZEA, β-Zearalenone, SEM, standard error of the mean.

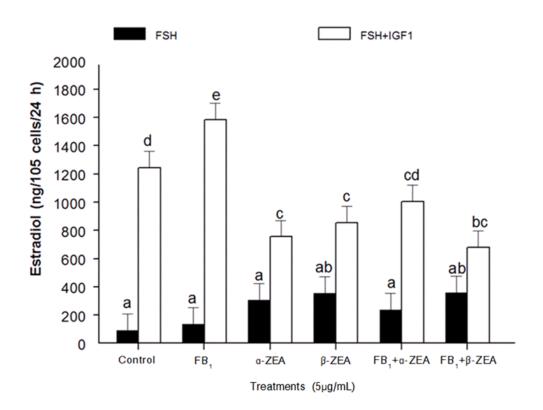


Figure 3.11: Interaction between FB<sub>1</sub>, α-ZEA or β-ZEA on estradiol production of bovine GC. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a-e) differ (P < 0.05). FB<sub>1</sub>, fumonisin B<sub>1</sub>; α-ZEA, α-Zearalenone;, β-ZEA, β-Zearalenone, SEM, standard error of the mean.

## 3.3.4 Experiment 4: Dose response of FB<sub>1</sub> and BEA on GC numbers and steroid production in the presence of FSH plus IGF1

Cell proliferation was not affected ( $P \ge 0.05$ ) by any dose of FB<sub>1</sub> (i.e., 0.3, 1, 3, 10  $\mu$ M), whereas BEA at the highest dose tested (10  $\mu$ M) decreased (72%; P < 0.01) cell numbers (Fig. 3.12). FB<sub>1</sub> at all doses did not significantly affect progesterone production, whereas BEA at 3 and 10  $\mu$ M strongly inhibited (P < 0.0001) progesterone production (Fig. 3.13). Estradiol production was inhibited (P < 0.05) by 25% and by 57% after exposure to FB<sub>1</sub> at 10  $\mu$ M and BEA at 3  $\mu$ M, respectively (Fig. 3.14).

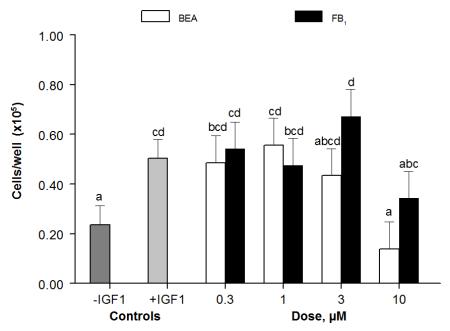
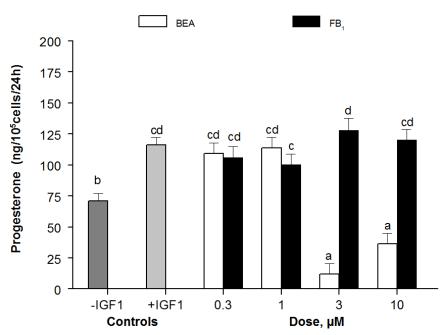


Figure 3.12: Effect of FB<sub>1</sub> and BEA on FSH plus IGF1-induced proliferation of bovine GC. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–d) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.



**Figure 3.13:** Effect of FB<sub>1</sub> and BEA on FSH plus IGF1-induced progesterone production by GC from bovine follicles. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–d) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.

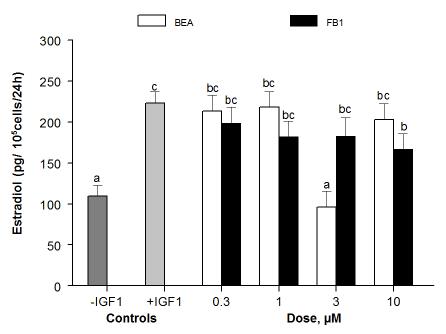
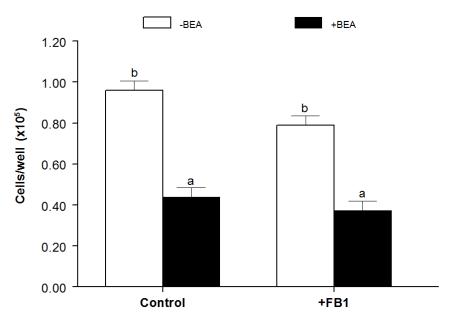


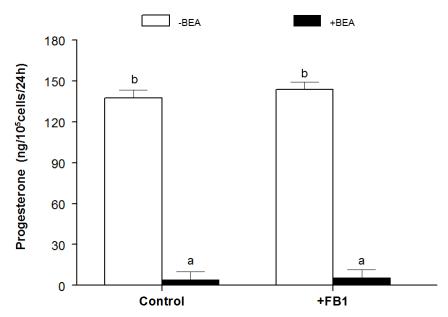
Figure 3.14: Effect of FB<sub>1</sub> and BEA on FSH plus IGF1-induced estradiol production by GC from bovine follicles. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–d) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.

# 3.3.5 Experiment 5: Individual and combined effects of FB<sub>1</sub> and BEA on GC numbers and steroid production in the presence of FSH plus IGF1

BEA (3  $\mu$ M) significantly inhibited (P< 0.0001) cell proliferation and progesterone and estradiol production by 54%, 97% and 80%, respectively (Fig. 3.15, 3.16, 3.17). FB<sub>1</sub> (3  $\mu$ M) did not affect (P > 0.05) any of the variables measured and did not influence the effect of BEA on cell proliferation (Fig. 3.15)or steroid production (Fig. 3.16, 3.17).



**Figure 3.15:** Interaction between FB<sub>1</sub> and BEA on FSH plus IGF1-induced proliferation of bovine GC. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–b) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.



**Figure 3.16:** Interaction between FB<sub>1</sub> and BEA on FSH plus IGF1-induced progesterone production by GC from bovine follicles. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–b) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.

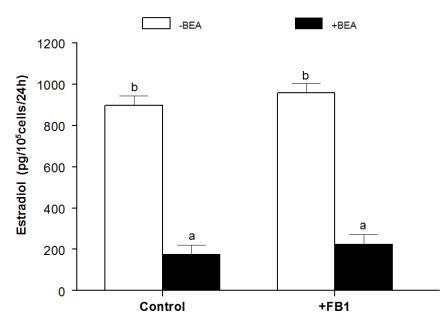


Figure 3.17: Interaction between FB<sub>1</sub> and BEA on FSH plus IGF1-induced estradiol production by GC from bovine follicles. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–b) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.

# 3.3.6 Experiment 6: Dose response of FB<sub>1</sub> with or without BEA on FSH plus IGF1-induced GC proliferation and steroid production

Cell proliferation was not affected (P  $\geq$  0.05) after exposure to FB<sub>1</sub> at 0.5, 1, 1.5 and 3  $\mu$ M alone or in combination with BEA (3  $\mu$ M), whereas at the highest dose tested (6  $\mu$ M) FB<sub>1</sub> in combination with BEA decreased (P < 0.05) cell numbers (Fig. 3.18). FB<sub>1</sub> alone had no significant effect (P > 0.05) on progesterone production (Fig. 3.19) whereas at 1 and 1.5  $\mu$ M FB<sub>1</sub> decreased (P < 0.05) estradiol production (Fig. 3.20). BEA at 3  $\mu$ M drastically inhibited both progesterone (88%; P < 0.001) and estradiol (96%; P < 0.0001) production, and the various doses of FB<sub>1</sub> did not influence these inhibitory effects (Fig. 3.19, 3.20).

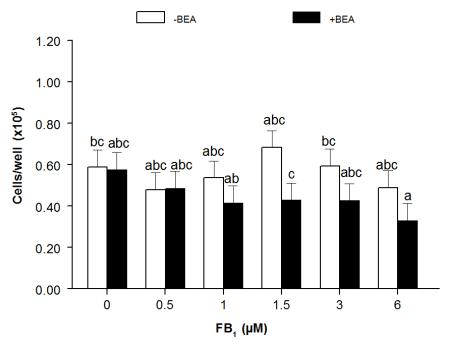
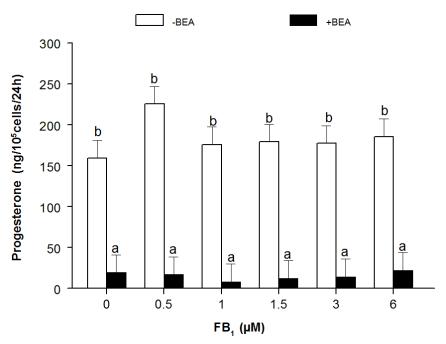
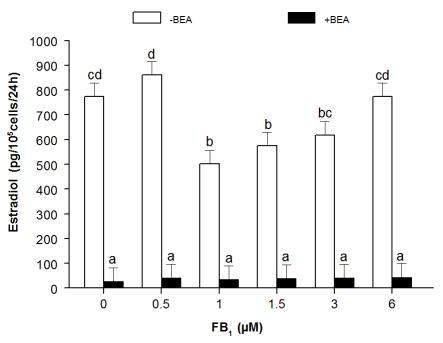


Figure 3.18: Effect of various doses of FB<sub>1</sub> with or without BEA on FSH plus IGF1-induced proliferation of bovine GC. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–c) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.



**Figure 3.19:** Effect of various doses of FB<sub>1</sub> with or without BEA on FSH plus IGF1-induced progesterone production by GC from bovine follicles. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–b) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.



**Figure 3.20:** Effect of various doses of FB<sub>1</sub> with or without BEA on FSH plus IGF1-induced estradiol production by GC from bovine follicles. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–b) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells

# 3.3.7 Experiment 7: Dose response of BEA with or without FB<sub>1</sub> on FSH plus IGF1-induced GC proliferation and steroid production

At the highest dose tested (6  $\mu$ M), BEA was found to inhibit (50%; P < 0.05) cell proliferation and no significant difference was detected in combination with FB<sub>1</sub> (3  $\mu$ M) (Fig. 3.21). BEA at 3 and 6  $\mu$ M was found to strongly decrease (P < 0.05) both progesterone and estradiol production, and FB<sub>1</sub> had little effect on these responses (Fig. 3.22, 3.23). FB<sub>1</sub> had no effect (P > 0.05) on progesterone production (Fig. 3.22) but at 3  $\mu$ M FB<sub>1</sub> decreased (P < 0.05) estradiol production alone and in combination with 1  $\mu$ M BEA (Fig. 3.23).

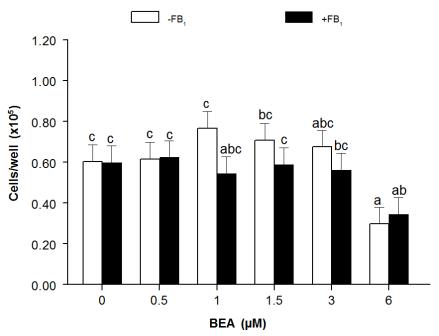


Figure 3.21: Effect of various doses of BEA with or without FB<sub>1</sub> on FSH plus IGF1-induced proliferation of bovine GC. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–c) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.

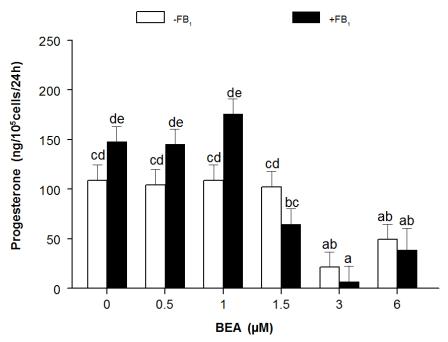
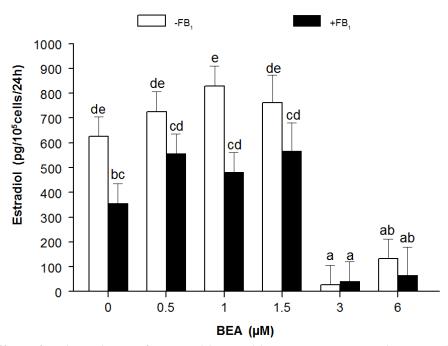


Figure 3.22: Effect of various doses of BEA with or without FB<sub>1</sub> on FSH plus IGF1-induced progesterone production by GC from bovine follicles. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–e) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.



**Figure 3.23:** Effect of various doses of BEA with or without FB<sub>1</sub> on FSH plus IGF1-induced estradiol production by GC from bovine follicles. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–e) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.

## 3.3.8 Experiment 8: Effect of FB1 and BEA on GC CYP11A1 and CYP19A1 mRNA

BEA (30  $\mu$ M) showed an inhibitory effect on FSH plus IGF1-induced CYP11A1 and CYP19A1 mRNA abundances (P < 0.05), whereas FB<sub>1</sub> (30  $\mu$ M) had no effect on gene expression (Fig. 3.24, 3.25).

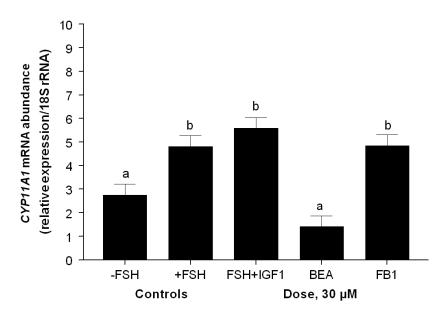


Figure 3.24: Effect of BEA and FB<sub>1</sub> on FSH plus IGF1-induced *CYP11A1* mRNA abundance in GC from bovine follicles. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–b) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.

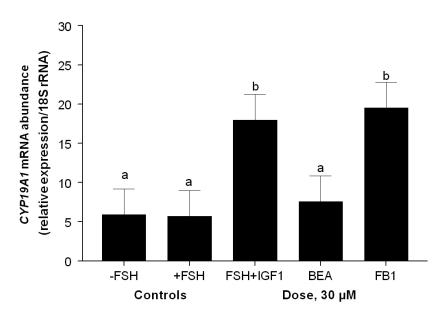
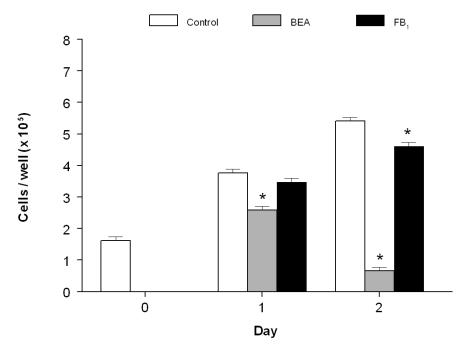


Figure 3.25: Effect of BEA and FB<sub>1</sub> on FSH plus IGF1-induced *CYP19A1* mRNA abundance in GC from bovine follicles. Values are means  $\pm$  SEM from three separate experiments. Means without a common letter (a–b) differ (P < 0.05). BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>; GC, granulosa cells.

## 3.3.9 Experiment 9: Effect of FB1 and BEA on serum-induced GC proliferation

Both FB<sub>1</sub> (10  $\mu$ M) and BEA (10  $\mu$ M) decreased (P < 0.001) GC proliferation induced by 10% FCS (Fig. 3.26). Cell numbers were decreased (P < 0.001) by 31% after 1 day of BEA treatment and by 88% and 15% after 2 days of BEA and FB<sub>1</sub> treatment, respectively (Fig. 3.26).



**Figure 3.26:** Effect of FB<sub>1</sub> and BEA on serum-induced proliferation of bovine GC. Within day of treatment, \* indicates mean differs (P < 0.001) from control value. Values are means  $\pm$  SEM from three separate experiments. BEA, beauvericin; FB<sub>1</sub>, fumonisin B<sub>1</sub>.

#### 3.4 Discussion

GC function is fundamental for normal folliculogenesis and oocyte growth and development (Spicer et al., 2001; Petro et al., 2012; Rawan et al., 2015). Previous *in vitro* studies with *Fusarium* mycotoxins such as DON and ZEA demonstrated that these mycotoxins are able to directly affect GC proliferation and steroid production and consequently the follicle development and oocyte function in different species, such as pigs (Ranzenigo et al., 2008; Caloni et al., 2009; Cortinovis et al., 2014) and cattle (Pizzo et al., 2016). Ruminants are considered more resistant to FB<sub>1</sub> than other species such as pigs and horses (Fink-Gremmels, 2008), but information on reproductive effects is lacking. The purpose of this study was to evaluate the *in vitro* toxicity of FB<sub>1</sub> using a bovine GC model.

Because the co-occurrence of Fusarium mycotoxins in commodities and their toxicological interaction have been demonstrated (Boutigny et al., 2012; Rodrigues and Naehrer, 2012), the combination of FB<sub>1</sub> with DON, α-ZEA, β-ZEA and BEA was also evaluated. In this study the results show that FB1 alone at all the concentrations tested (0.042-10 µM) did not affect IGF1-induced GC proliferation. Differently, FB<sub>1</sub> at 30 ng/mL (0.042  $\mu M)$  in combination with  $\beta\text{-ZEA}$  at 30 ng/mL (0.094  $\mu M)$  (Fig. 3.3), had a stimulatory effect on GC numbers in the presence of IGF1, while β-ZEA alone at 30 ng/mL (0.094 μM) had no effect on cell proliferation (Fig. 3.3). FB<sub>1</sub> at 5 μg/mL (6.9 μM) in combination with β-ZEA at 5 μg/mL (15.6 μM) inhibited IGF1-induced cell proliferation (Fig. 3.9), but this effect was also observed with β-ZEA alone (15.6 μM) (Fig. 3.9), and is in agreement with a previous study conducted on porcine GC (Tiemann et al., 2003). In accordance with previous studies (Ranzenigo et al., 2008; Pizzo et al., 2016), α-ZEA and FB<sub>1</sub> alone had not effect on GC proliferation, while co-exposure of FB<sub>1</sub> (6.9  $\mu$ M) and  $\alpha$ -ZEA (15.6  $\mu$ M) decreased IGF1-induced cell proliferation (Fig. 3.9), indicating that these two mycotoxins may interact inducing an inhibition of cell proliferation.

Exposure of bovine GC to DON alone or combined with FB<sub>1</sub> had no effect on cell proliferation, whereas previous studies conducted on porcine GC reported conflicting results on DON activity, showing either an inhibitory effect (Ranzenigo et al., 2008) or a stimulatory effect (Medvedova et al., 2011) depending on the dose of DON and the presence of  $\alpha$ -ZEA. All these results suggest that the effects of FB<sub>1</sub>, DON,  $\beta$ -ZEA and  $\alpha$ -ZEA on GC proliferation may be species-specific and influenced by their interaction. In the present study, the effect of BEA on GC proliferation was investigated for the first time. At concentrations  $\geq$  6  $\mu$ M (Fig. 3.21), BEA was found to inhibit GC proliferation confirming results previously reported for other cell types (Jestoi, 2008). According to previous studies (Klarić et al., 2008; Ferrer et al., 2009; Ruiz et al., 2011; Prosperini et

al., 2012; Mallebrera et al., 2016) BEA exhibits potent cytotoxicity against several mammalian cell lines. In details, BEA was found to reduce cell viability in a time and concentration-dependent manner in Vero cells, CHO-K1 cells and Caco-2 cells at 6.25 to 10.02 (Ruiz et al., 2011), 5  $\mu$ M (Mallebrera et al., 2016) and at 3.125 to 25  $\mu$ M (Prosperini et al., 2012), respectively.

The present study evaluated also the effects of Fusarium mycotoxins on GC steroidogenesis. FB<sub>1</sub> alone at all the concentrations tested did not affect GC P4 production. On the contrary, when combined with  $\beta$ -ZEA at 30 ng/mL (0.094  $\mu$ M) and DON at 100 ng/mL (0.33  $\mu$ M) plus  $\beta$ -ZEA at 30 ng/mL (0.094  $\mu$ M), FB<sub>1</sub> at 100 ng/mL (0.13  $\mu$ M) stimulated and inhibited P4 production, respectively.

The lack of an effect of FB<sub>1</sub> on P4 production is also supported by the results showing no effect of FB<sub>1</sub> on CYP11A1 mRNA abundance in GC. Differently BEA was found to inhibit P4 production and this effect was associated with a significant suppression of CYP11A1 mRNA abundance, suggesting that BEA may alter P4 production via a change in CYP11A1 mRNA abundance.

As regards E2 production, in this study FB<sub>1</sub> at 30 (0.042  $\mu$ M) and 100 ng/mL (0.13  $\mu$ M) did not affect E2 production (Fig. 3.5), but at 5 µg/mL (6.9 µM) increased E2 production (Fig. 3.11), whereas, at the same concentration (i.e., 5 µg/mL corresponding to 6.9  $\mu$ M ) both  $\alpha$ -ZEA and  $\beta$ -ZEA alone inhibited E2 production (Fig. 3.11). The combination of FB<sub>1</sub> at 5  $\mu$ g/mL (6.9  $\mu$ M) with  $\alpha$ -ZEA at 5  $\mu$ g/mL (15.6  $\mu$ M) and  $\beta$ -ZEA at 5 µg/mL (15.6 µM) confirms the inhibition of E2 production induced by these two ZEA metabolites (Fig. 3.11). Indeed, FB<sub>1</sub> combined with  $\alpha$ -ZEA at 5  $\mu$ g/mL (15.6  $\mu$ M) had no effect on E2 production, whereas in combination with  $\beta$ -ZEA at 5  $\mu$ g/mL (15.6) μM) an inhibitory effect on E2 release was observed. It is well established that estrogenic α-ZEA and β-ZEA effects are mediated by binding to intracellular estrogen receptors (Parveen et al., 2009). The inhibition on E2 production of ZEA major metabolites, clearly highlighted as a result of co-exposure with FB<sub>1</sub>, suggests an interference on aromatase activity. FB<sub>1</sub> at 30 ng/ml (0.042 μM) and 100 ng/mL (0.13 μM) either with or without β-ZEA had no effect on the inhibition of E2 production induced by DON at 100 ng/mL (0.33 μM) (Fig. 3.8), an effect previously reported (Pizzo et al., 2016). The BEA-induced marked inhibition of E2 production by GC was associated with a significant decrease in CYP19A1 mRNA abundance, suggesting that BEA may alter E2 production via suppression of CYP19A1 mRNA abundance. Previous studies demonstrated a role of CART (cocaine and amphetamine regulated transcript system) system on the regulation of bovine GC E2 production (Kobayashi et al., 2006; Lv et al., 2009). This effect was linked to the inhibition induced by CART on several components of the FSH signal transduction pathway (Lv et al., 2009) that, since FSH stimulates E2 production, may explain the negative effects of CART observed on bovine GC (Lv et al., 2009). This mechanism of action could be involved for the inhibitory effects of BEA on

E2 production and GC CYP19A1 mRNA levels observed in the present study. BEA was found to reversibly inhibit L-type calcium channels in a dose-dependent manner in a neuronal cell line (Wu, 2002) and FSH-induced increase in GC calcium uptake is mediated via inhibition of L-type calcium channel activity (Peters et al., 2004; Kobayashi et al., 2006), but whether blockade of calcium channels is involved in BEA-induced inhibition of E2 production will require further study.

The present study has been carried out with FB<sub>1</sub> alone and in co-exposure firstly with DON and ZEA metabolites ( $\alpha$ -ZEA and  $\beta$ -ZEA) and secondly with BEA, an emerging mycotoxin, demonstrating effects on cell proliferation and steroidogenesis using a bovine GC *in vitro* model, suggesting possible reproductive effects of these mycotoxins in cattle.

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4. Effects of fumonisin B <sub>1</sub> alone and combined w	ith
beauvericin on Caco-2 in vitro intestinal model	

## 4.1 Introduction

Mycotoxins are secondary metabolites produced by molds, such as *Aspergillus*, *Fusarium* and *Penicillium* (Osweiler, 2000). *Fusarium* mycotoxins are world-spread mycotoxins naturally occurring in cereals (Jurjevic, 2002; Logrieco et al., 2003; Domijan et al., 2005; Molinié et al., 2005; Ferrigo et al., 2016) and animal and human exposure is well established (Diekman and Green 1992; Binder et al., 2007; Fink-Gremmels, 1999; Prosperini, et al., 2012; Wu et al., 2014; EFSA 2014).

Fumonisins are a group of mycotoxins mainly produced by Fusarium verticillioides, Fusarium proliferatum, Fusarium napiforme and Fusarium nygamai that have been shown to occur worldwide primarily in corn (Glenn, 2007; Voss et al., 2007).

The most important group of fumonisins is the B series including fumonisin B<sub>1</sub> (FB<sub>1</sub>), B<sub>2</sub> (FB<sub>2</sub>) and B<sub>3</sub> (FB<sub>3</sub>) (EFSA 2005). In 2002 the IARC classified FB<sub>1</sub> as possibly carcinogenic to humans (group 2B) (IARC, 2002; Santini et al., 2015; Hove et al., 2016) and several studies reported that FB<sub>1</sub> is associated with an increased prevalence of esophageal cancer in humans (Marasas et al., 1988; Sydenham et al. 1990; Rheder et al., 1992; Hove et al., 2016).

The complex mechanism of action of FB<sub>1</sub> is based on the inhibition of sphingosine (sphinganine) N-acetyltransferase (ceramide synthase) (Wang et al., 1991; Voss et al., 2002, 2007; Luongo et al. 2008; Gbore et al., 2012) with the consequent alteration of the sphingolipid metabolism (Wang et al., 1991; Wang et al; 1992; Merrill Jr. et al., 2001; Marasas et al., 2004; Voss et al., 2007; Smith, 2012). Specifically FB<sub>1</sub>, structurally similar to sphingoid bases, inhibits ceramide synthase and thus disrupts the *de novo* biosynthesis of ceramide and sphingolipid metabolism (Wang et al., 1991). FB<sub>1</sub> is a competitive inhibitor with respect to both substrates of ceramide synthase (Merril Jr. et al., 2001; Marasas et al., 2004; Voss et al., 2007). This leads to the blockage of complex sphingolipid biosynthesis, essential for cell regulation, and to the accumulation of sphinganine, and, to a lesser degree, sphingosine (Wang et al., 1992; Merril Jr. et al., 2001; Marasas et al., 2004; Smith, 2012).

Fumonisins have been found to commonly occur in cereal grains and animal feed in combination with other *Fusarium* mycotoxins including beauvericin (BEA), a so-called emerging mycotoxin (Jestoi, 2008; EFSA, 2014).

BEA is a ionophoric molecule that can form lipophilic complexes with cations and transport them into biological membranes altering cell homeostasis (Hilgenfeld and Saenger, 1982; Jestoi et al., 2008; Prosperini et al., 2012; Fernandez-Blanco et al., 2016). In fact, it is well-known that an increase in the intracellular concentration of cations such as calcium and the subsequent activation of calcium-dependent endonucleases lead to DNA fragmentation which is involved in apoptosis (Speijers and Speijers, 2004; Kouadio et al., 2007).

Recent studies reported the *in vitro* adverse effects of BEA on human, porcine and rodent cell lines (Jestoi, 2008; Prosperini et al., 2012; Fernandez Blanco et al., 2016; Mallebrera et al., 2016). Schoevers et al. (2016) studied the effects of BEA on porcine cumulus cells, oocytes and embryos. In this study the cumulus-oocyte-complexes and developing embryos were exposed to BEA at concentrations ranging from 0.31 to 10  $\mu$ M and the effects of this mycotoxin on viability, progesterone synthesis and apoptosis were evaluated (Schoevers et al., 2016). Schoevers et al. (2016) demonstrated that BEA reduced the developmental competence of both the maturing oocytes and the two-four cell stage embryos in pigs, and that BEA only affected the rate of developing embryos at concentrations exceeding 0.5  $\mu$ M. BEA toxic effects on human cell lines were investigated by Prosperini et al. (2012). The authors evaluated the cytotoxicity of BEA on human Caco-2 cells testing concentrations ranging from 0.6 to 30  $\mu$ M. The IC50 obtained for BEA was 20.62  $\pm$  6.9  $\mu$ M at 24 h of exposure and 12.75  $\pm$  4.8 at 48 h (Prosperini et al., 2012).

Recently, exposure of Caco-2 cells to  $0.001-10~\mu M$  BEA showed that BEA at 1  $\mu M$  and  $10~\mu M$  had cytotoxic effects on Caco-2 cells (Fernandez-Blanco et al., 2016).

The *in vitro* absorption and toxicity of FB<sub>1</sub> have been investigated with different intestinal models (Caloni et al., 2002; Bouhet et al., 2004; De Angelis et al., 2005; Loiseau et al., 2007; Ulluwishewa et al., 2011; Minervini et al., 2014; Romero et al., 2016; Wentzel et al., 2016). *In vitro* studies with Caco-2 cell monolayers (Caloni et al., 2002) demonstrated low toxicity of FB<sub>1</sub> and its metabolites after 48 h of exposure to concentrations ranging from 1 to 138 μM, even though FB<sub>1</sub> appeared to be the most active. The results of a subsequent study carried out with Caco-2 cells cultured on inserts (Caloni et al., 2005) found that FB<sub>1</sub> was not absorbed by Caco-2 cells and did not affect the barrier integrity. Recently, exposure of Caco-2 cells to 0.001–10 μM FB<sub>1</sub> revealed no cytotoxic effect of FB<sub>1</sub> at all the concentrations tested (Fernandez-Blanco et al., 2016).

In IPEC-1, a porcine cell line, a significant TEER decrease was reported after 13 days of exposure to FB<sub>1</sub> (50  $\mu$ M) (Bouhet et al., 2004).

The aim of this study was to clarify the *in vitro* effects of FB<sub>1</sub> and BEA, alone and combined, on human intestinal Caco-2 cells cultured on inserts through the evaluation of TEER and cytokine release.

## 4.2 Materials and methods

#### 4.2.1 Chemicals

Dulbecco's Modified Eagles Medium (DMEM) high glucose, heat inactivated fetal bovine serum (FBS), glutamine, non-essential amino acids (NEAA), N-2-hydroxyethylpiperazine-N-2'-ethanesulfonic acid (HEPES), penicillin/streptomycin were all purchased from Gibco (Thermo Fisher Scientific, Inc., Waltham, MA, USA). Fumonisin B<sub>1</sub>, beauvericin and methanol were all obtained from Sigma-Aldrich Chemical Company (St. Louis, MO, USA).

## 4.2.2 Cell culture

Caco-2 cells were purchased from Sigma-Aldrich Chemical Company (St. Louis, MO, USA) and routinely grown in an humidified atmosphere of 5% carbon dioxide at 37°C in DMEM high glucose standard medium supplemented with 100 IU/ml penicillin and 100 μg/ml streptomycin, 4 mmol/l glutamine, 1% NEAA, 10 mM HEPES and 10% heat inactivated FBS. Starting from a cell suspension in culture medium (3 × 10<sup>5</sup> cells/ml), the cells were seeded at a density of 1.5 × 10<sup>5</sup> cells/filter on 4 μm pore size 12-well plate polyethylene terephthalate (PET) transparent membrane inserts (Millicell®, Millipore Corporation) (Fig. 4.1); 0.5 ml of cellular suspension were added in the apical (Ap) compartment and 1.5 ml of supplemented DMEM in the basolateral (Bl) compartment of each insert. The plate was then shaken gently to avoid non uniform cell distribution.

The plates were transferred to a 37°C, 5% CO<sub>2</sub> incubator. Cells were allowed to differentiate for 21 days with regular medium changes three times per week.

The following procedure was used to change the medium on the inserts: first the medium from the Bl side of a single well was removed, then the medium from the Ap side of the same well was slowly removed avoiding to touch the cellular monolayer with the pipet tip; the medium in the Ap compartment was replaced (0.5 ml) first, then in the Bl compartment (1.5 ml). This procedure was repeated for each well. The culture was regularly checked with an inverted microscope to identify contaminations and/or morphological variations. Treatments were applied at the end of the differentiation process.



Figure 4.1: Caco-2 cells cultured on 12 well plate inserts

## 4.2.3 Barrier Integrity Assessment (Trans Epithelial Electrical Resistance Evaluation)

Barrier integrity was assessed by measuring the trans-epithelial electrical resistance (TEER) which quantifies the movement of ions across the cell barrier. TEER values were recorded in the culture medium at 37°C using an epithelial volthommeter (Millicell-ERS, Millipore; Fig: 24) just before (0 h) and 1, 2 and 24 h after treatment.

Before the measurement, the chop-stick electrodes were sterilized. After sterilization, the electrodes were placed in the culture medium, both in the Ap and Bl compartments. Three separate measurements were quickly performed for each filter and the TEER values were calculated with the following equation:

TEER =  $(\Omega \text{ cell monolayer} - \Omega \text{ filter cell-free}) x \text{ filter area}$ 

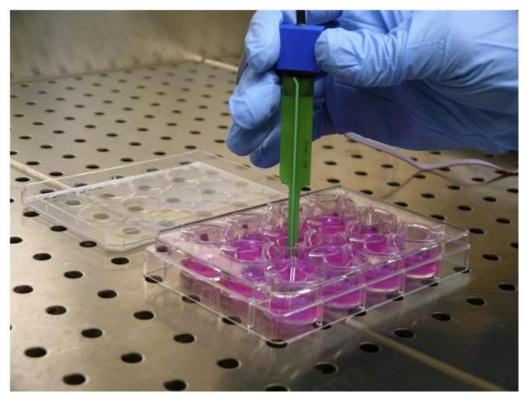


Figure 4.2: measurement of TEER

## 4.2.4 Measurement of pro-inflammatory mediator release

After 24 hours of treatment, culture medium was collected from individual wells and frozen at -80°C for subsequent pro-inflammatory cytokine release determination. In all experiments, the culture medium was analysed for the presence of interleukin-6 (IL-6), interleukin-8 (IL-8) and tumour necrosis factor-α (TNF-α) using commercially available quantitative ELISA assay kits (Sigma-Aldrich Chemical Company, St. Louis, MO, USA).

## 4.2.4.1 IL-6 determination

The ELISA procedure was performed according to the manufacturer's instructions of the Human IL-6 ELISA Kit (Sigma-Aldrich Chemical Company, St. Louis, MO, USA).

Briefly, all reagents and samples were taken to room temperature (18-25°C) before use. 100 µl of each standard and sample were placed into appropriate wells, covered and incubated for 2.5 h at room temperature. All standards were performed in duplicate.

After incubation, the solution was discarded and the wells were washed 4 times with 1x Wash Buffer (300 µl). After the washing step, 100 µl of 1x prepared Biotinylated

Detection Antibody were introduced in each well and the plate was incubated for 1 h at room temperature. The solution was then discarded and the washing step repeated. Successively,  $100~\mu l$  of prepared HRP-Streptavidin Solution were added to each well and incubation was carried out for 45 minutes. The solution was then discarded and after the washing step,  $100~\mu l$  of ELISA Colorimetric TMB Reagent were added to each well. After 30 minutes of incubation,  $50~\mu l$  of Stop Solution were introduced in each well and the optical density (OD) was immediately measured at 450 nm by an ELISA reader (Multiskan GO microplate spectrophotometer, Thermo Scientific, Waltham, MA, USA). A calibration curve was constructed with eight standard concentrations: 0~pg/ml - 1.37~pg/ml - 4.12~pg/ml - 12.35~pg/ml - 37.04~pg/ml - 111.1~pg/ml - 333.3~pg/ml - 1000~pg/ml. After the subtraction of the average zero standard OD, IL-6 sample concentration was measured by interpolation from the calibration curve.

## 4.2.4.2 IL-8 determination

The ELISA procedure was performed according to the manufacturer's instructions of the Human IL-8 / CXCL8 ELISA Kit (Sigma-Aldrich Chemical Company, St. Louis, MO, USA).

Briefly, all reagents and samples were taken to room temperature (18-25°C) before use. 100 µl of each standard and sample were placed into appropriate wells, covered and incubated for 2.5 h at room temperature.

All standards were performed in duplicate and samples were diluted (1:3).

After incubation, the solution was discarded and the wells were washed 4 times with 1x Wash Buffer (300 µl). After the washing step, 100 µl of 1x prepared Biotinylated Detection Antibody were introduced in each well and the plate was incubated for 1 h at room temperature. The solution was then discarded and the washing step repeated. Successively, 100 µl of prepared HRP-Streptavidin Solution were added to each well and incubation was carried out for 45 minutes. The solution was then discarded and after the washing step, 100 µl of ELISA Colorimetric TMB Reagent were added to each well. After 30 minutes of incubation, 50 µl of Stop Solution were introduced in each well and the optical density (OD) was immediately measured at 450 nm by an ELISA reader (Multiskan GO microplate spectrophotometer, Thermo Scientific, Waltham, MA, USA). A calibration curve was constructed with eight standard concentrations: 0 pg/ml – 0.8 pg/ml – 2.5 pg/ml – 7.4 pg/ml – 22.2 pg/ml – 66.7 pg/ml – 200 pg/ml – 600 pg/ml. After the subtraction of the average zero standard OD, IL-8 sample concentration was measured by interpolation from the calibration curve.

## 4.2.4.3 Tumor Necrosis Factor α determination

The ELISA procedure was performed according to the manufacturer's instructions of the Human Tumor Necrosis Factor  $\alpha$  ELISA Kit (Sigma-Aldrich Chemical Company, St. Louis, MO, USA).

Briefly, all reagents and samples were taken to room temperature (18-25°C) before use. 100 µl of each standard and sample were placed into appropriate wells, covered and incubated for 2.5 h at room temperature. All standards were performed in duplicate. After incubation, the solution was discarded and the wells were washed 4 times with 1x Wash Buffer (300 µl). After the washing step, 100 µl of 1x prepared Biotinylated Detection Antibody were introduced in each well and the plate was incubated for 1 h at room temperature. The solution was then discarded and the washing step repeated. Successively, 100 µl of prepared HRP-Streptavidin Solution were added to each well and incubation was carried out for 45 minutes. The solution was then discarded and after the washing step, 100 µl of ELISA Colorimetric TMB Reagent were added to each well. After 30 minutes of incubation, 50 µl of Stop Solution were introduced in each well and the optical density (OD) was immediately measured at 450 nm by an ELISA reader (Multiskan GO microplate spectrophotometer, Thermo Scientific, Waltham, MA, USA). A calibration curve was constructed with eight standard concentrations: 0 pg/ml - 24.58pg/ml - 61.44 pg/ml - 153.6 pg/ml - 384 g/ml - 960 pg/ml - 2400 pg/ml - 6000pg/ml. After the subtraction of the average zero standard OD, TNF-α sample concentration was measured by interpolation from the calibration curve.

## 4.2.5 Experimental Design

Experiment 1 was performed to evaluate the dose response of fumonisin  $B_1$  (FB<sub>1</sub>) on Caco-2 barrier integrity and the release of the pro-inflammatory cytokines IL-6, IL-8 and TNF- $\alpha$ . Caco-2 cells were cultured as previously described and, after differentiation, cells were treated with different concentrations of FB<sub>1</sub> (0, 0.5, 1.5, 3  $\mu$ M) in triplicate from both Ap and Bl sides. The barrier integrity was evaluated after 1, 2 and 24 h of exposure to FB<sub>1</sub> by measuring TEER. After 24 h of treatment, medium was collected for IL-6, IL-8 and TNF- $\alpha$  determination.

Experiment 2 was performed to evaluate the dose response of beauvericin (BEA) on Caco-2 barrier integrity and the release of the pro-inflammatory cytokines IL-6, IL-8 and TNF- $\alpha$ . Caco-2 cells were cultured as previously described and, after differentiation, cells were treated with different concentrations of BEA (0, 0.5, 1.5, 3, 6  $\mu$ M) in triplicate from both Ap and Bl sides. The barrier integrity was evaluated after 1, 2 and 24 h of

exposure to BEA by measuring TEER. After 24 h of treatment, medium was collected for IL-6, IL-8 and TNF-α determination.

Experiment 3 was designed to evaluate the effects of FB<sub>1</sub> (1.5  $\mu$ M) and BEA (3  $\mu$ M), alone and combined, on Caco-2 barrier integrity and the release of the pro-inflammatory cytokines IL-6, IL-8 and TNF- $\alpha$ . Caco-2 cells were cultured as previously described and, after differentiation, cells were treated with FB<sub>1</sub> (0 or 1.5  $\mu$ M) and BEA (0 or 3  $\mu$ M) in triplicate from both Ap and Bl sides. The barrier integrity was evaluated after 1, 2 and 24 h of exposure to treatments by measuring TEER. After 24 h of treatment, medium was collected for IL-6, IL-8 and TNF- $\alpha$  determination.

## 4.2.6 Statistical analysis

Each experiment was performed in triplicate. Results are expressed as mean  $\pm$  standard deviations (SD). Statistical evaluation was performed by two tailed Student's t-test. The level of significance was established at P < 0.05.

## 4.3 Results

## 4.3.1 Experiment 1: effect of fumonisin $B_1$ (FB<sub>1</sub>) on Caco-2 barrier integrity and cytokine release

Caco-2 cells were treated with different concentrations of FB<sub>1</sub> (0, 0.5, 1.5, 3  $\mu$ M) from both Ap and Bl sides. The barrier integrity was evaluated after 1, 2 and 24 h of exposure to FB<sub>1</sub> by measuring TEER. After 24 h of treatment, medium was collected for IL-6, IL-8 and TNF- $\alpha$  determination.

No significant (P  $\geq$  0.05) effect on TEER was observed after 24 h of both Ap and Bl exposure to all doses (0.5, 1.5, 3  $\mu$ M) of FB<sub>1</sub> (Fig. 4.3 and 4.4). On one hand, no significant release of the inflammatory mediators IL-6 and TNF- $\alpha$  was observed after Ap and Bl exposure to FB<sub>1</sub> (Fig. 4.5 and 4.7) at all doses (0.5, 1.5, 3  $\mu$ M). On the other hand, a significant (P < 0.05) increase of IL-8 release was induced by Bl exposure to FB<sub>1</sub> at 3  $\mu$ M (Fig. 4.6).

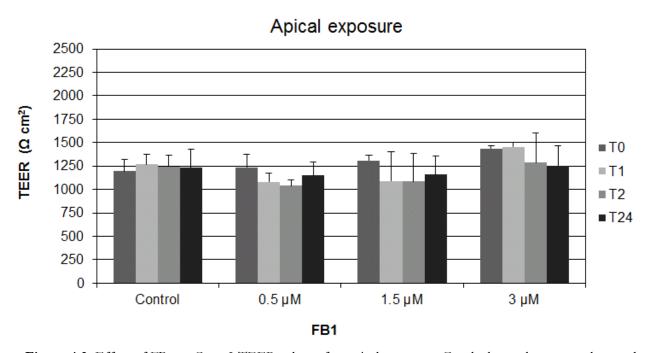
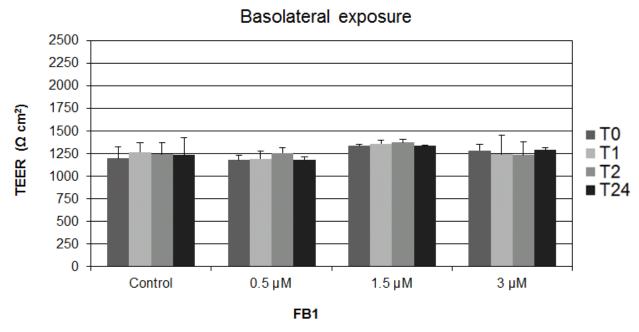


Figure 4.3: Effect of FB<sub>1</sub> on Caco-2 TEER values after apical exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.



**Figure 4.4:** Effect of FB<sub>1</sub> on Caco-2 TEER values after basolateral exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.

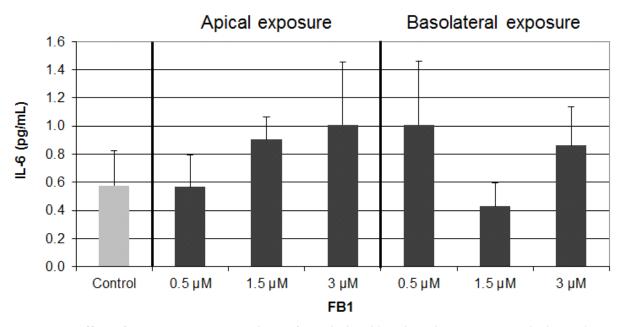
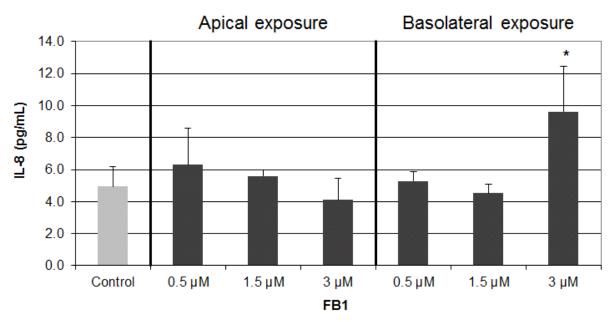
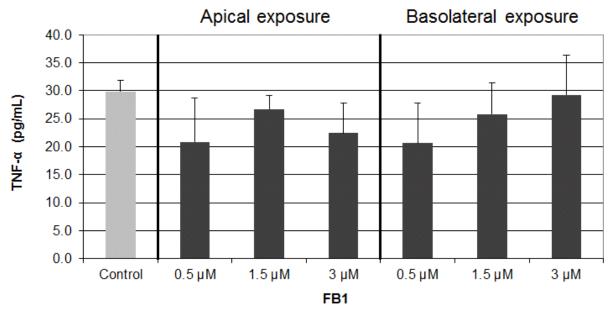


Figure 4.5: Effect of FB<sub>1</sub> on Caco-2 IL-6 release after apical and basolateral exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.



**Figure 4.6:** Effect of FB<sub>1</sub> on Caco-2 IL-8 release after apical and basolateral exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.



**Figure 4.7:** Effect of FB<sub>1</sub> on Caco-2 TNF- $\alpha$  release after apical and basolateral exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.

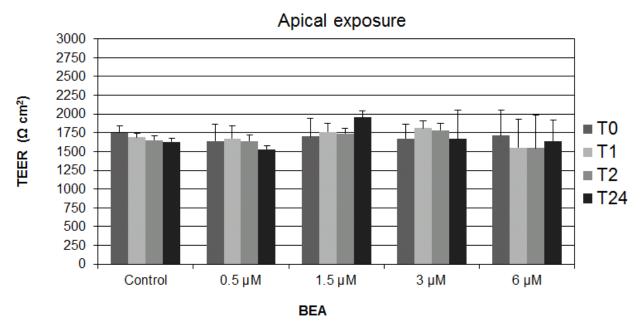
## 4.3.2 Experiment 2: effect of beauvericin (BEA) on Caco-2 barrier integrity and cytokine release

Caco-2 cells were treated with different concentrations of BEA (0, 0.5, 1.5, 3, 6  $\mu$ M) from both Ap and Bl sides. The barrier integrity was evaluated after 1, 2 and 24 h of exposure to BEA by measuring TEER. After 24 h of treatment, medium was collected for IL-6, IL-8 and TNF- $\alpha$  determination.

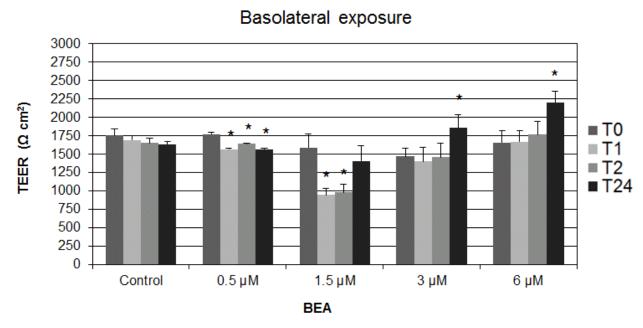
TEER was not significantly affected ( $P \ge 0.05$ ) by Ap exposure to all doses of BEA (Fig. 4.8). On the contrary, a significant (P < 0.05) TEER decrease was observed after Bl exposure to BEA at 0.5 and 1.5  $\mu$ M starting from the first hour of treatment (Fig. 4.9). This decrease in TEER was no more observed after 24 h of Bl exposure to BEA at 1.5  $\mu$ M. Differently, at higher concentrations (3 and 6  $\mu$ M), BEA was found to significantly (P < 0.05) increase TEER after 24 h of Bl exposure (Fig. 4.9).

A significant release of the inflammatory mediator IL-6 was observed after Ap exposure to BEA at 3 and 6  $\mu$ M and after Bl exposure to BEA at 1.5, 3 and 6  $\mu$ M (Fig. 4.10). Regarding IL-8, a significant (P < 0.05) release was induced by Ap exposure to BEA at 3 and 6  $\mu$ M and by Bl exposure to BEA only at 6  $\mu$ M (Fig. 4.11).

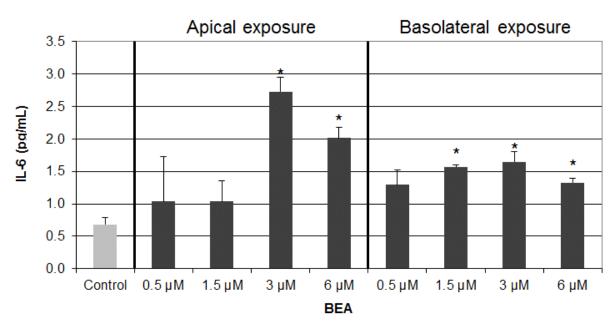
Concerning TNF- $\alpha$ , a significant release was observed after Ap exposure to 0.5 and 1.5  $\mu$ M BEA and after Bl exposure to 1.5  $\mu$ M BEA (Fig. 4.12).



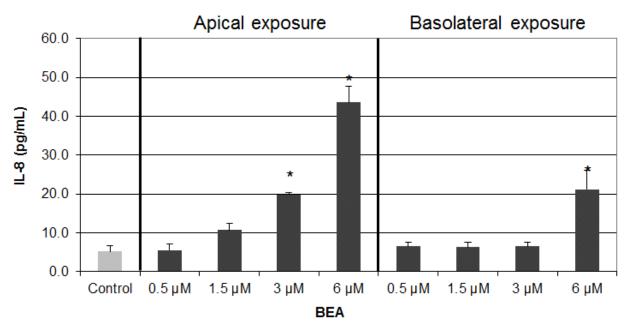
**Figure 4.8:** Effect of BEA on Caco-2 TEER values after apical exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.



**Figure 4.9:** Effect of BEA on Caco-2 TEER values after basolateral exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.



**Figure 4.10:** Effect of BEA on Caco-2 IL-6 release after apical and basolateral exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.



**Figure 4.11:** Effect of BEA on Caco-2 IL-8 release after apical and basolateral exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.

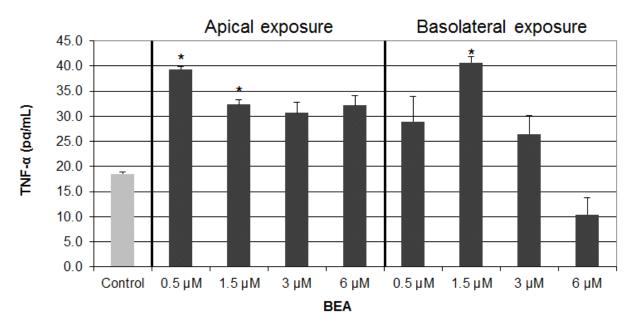


Figure 4.12: Effect of BEA on Caco-2 TNF- $\alpha$  release after apical and basolateral exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.

# 4.3.3 Experiment 3: individual and combined effects of fumonisin $B_1$ and beauvericin (BEA) on Caco-2 barrier integrity and cytokine release

Caco-2 cells were treated with FB<sub>1</sub> (0 or 1.5  $\mu$ M) and BEA (0 or 3  $\mu$ M) from both Ap and Bl sides. The barrier integrity was evaluated after 1, 2 and 24 h of exposure to treatments by measuring TEER. After 24 h of treatment, medium was collected for IL-6, IL-8 and TNF- $\alpha$  determination.

No significant ( $P \ge 0.05$ ) effect was observed on TEER after 24 h of Ap exposure to FB<sub>1</sub> at 1.5  $\mu$ M, BEA at 3  $\mu$ M and the combination of these mycotoxins (Fig. 4.13). Only after 24 h of Bl exposure to BEA at 3  $\mu$ M a significant (P < 0.05) increase of TEER was observed (Fig. 4.14). This TEER increase was not observed after Bl exposure to BEA at 3  $\mu$ M in combination with FB<sub>1</sub> at 1.5  $\mu$ M (Fig. 4.14)

FB<sub>1</sub> and BEA, alone and combined, had no significant effects on cytokine release (Fig. 4.15, 4.16, 4.17).

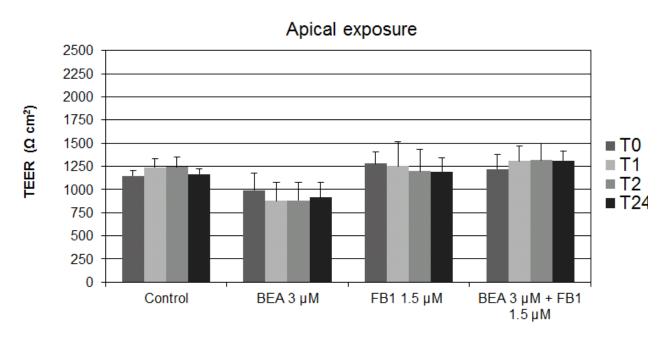


Figure 4.13: Effect of FB<sub>1</sub> with or without BEA on Caco-2 TEER values after apical exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.

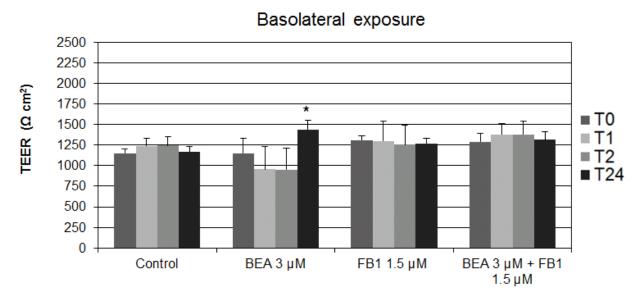


Figure 4.14: Effect of FB<sub>1</sub> with or without BEA on Caco-2 TEER values after basolateral exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.

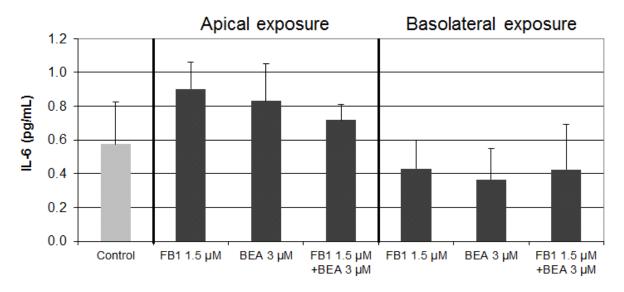


Figure 4.15: Effect of FB<sub>1</sub> with or without BEA on Caco-2 IL-6 release after apical and basolateral exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.

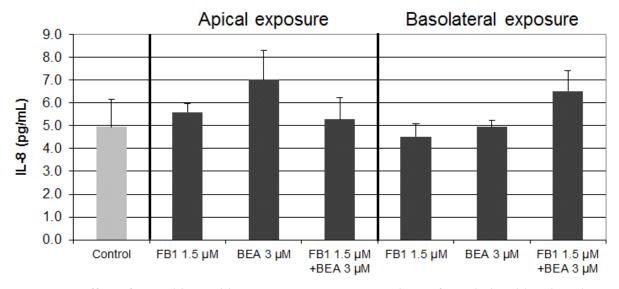


Figure 4.16: Effect of FB<sub>1</sub> with or without BEA on Caco-2 IL-8 release after apical and basolateral exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.

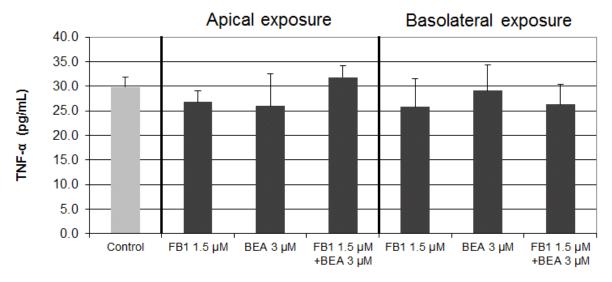


Figure 4.17: Effect of FB<sub>1</sub> with or without BEA on Caco-2 TNF- $\alpha$  release after apical and basolateral exposure. Graph shows the mean values and standard deviations (n=3). (\*) P < 0.05.

## 4.4 Discussion

The gastrointestinal tract is an important environment responsible for absorbing nutrients, protecting the body against pathogens entering with food and preventing the loss of important compounds such as water and solutes (Gordon et al., 2015). The integrity of the intestinal barrier is crucial for the maintenance of the homeostasis (Goto and Kiyono, 2012). An impairment of the intestinal barrier function may cause autoimmune responses, inflammation and atopic diseases (Antonissen et al., 2014). The trans-epithelial electrical resistance (TEER) is a parameter that gives information on the barrier integrity and its decrease corresponds to an impairment (Marin et al., 2015). TEER is considered a good indicator of the organization of the TJ proteins (Pinton et al., 2009). TJ proteins seal the intercellular space between adherent epithelial cells, thus preventing paracelluar transport of luminal antigens (Schneeberger and Lynch, 1992; Liu et al., 2000; Tsukita et al., 2001; Maresca and Fantini, 2010). TJ proteins are therefore involved in barrier function and their damage leads to an increase of the intestinal permeability and thus to an increased transepithelial passage of bacteria and antigens (Maresca and Fantini, 2010). The number of TJ strands and their ramification depend on the cell type, producing marked variation in the morphology of TJ strand networks. The main TJ integral membrane proteins are occludins (OCLN), claudins (CLDN) and junctional adhesion molecules (JAM) (Tsukita et al., 2001; Maresca and Fantini, 2010). The ability of FB<sub>1</sub> to cause toxicity on intestinal epithelial cell lines have been already reported. Bouhet et al. (2004) showed that FB<sub>1</sub> at concentrations ranging from 2 to 700 μM decreased in a time- and dose-dependent manner TEER of IPEC-1 cells (Bouhet et al., 2004). The authors reported a significant TEER decrease after 13-day exposure to FB<sub>1</sub> at 50 μM and a drastic impairment of the intestinal barrier integrity after 15 to 18 days of exposure to high concentrations of FB<sub>1</sub> (200 and 500 μM) (Bouhet et al., 2004). FB<sub>1</sub> (1 to 138 μM) was previously reported to not exert cytotoxic effects on Caco-2 cells (Caloni et al., 2002). Only a transient decrease of TEER was observed after 6-h exposure to the highest concentration tested, without FB<sub>1</sub> epithelial passage (Caloni et al., 2005; De Angelis et al., 2005). Recently, after treatments with FB<sub>1</sub> at 1, 3, 10, 30 µM a reduction in mRNA levels of CLDN3 and CLDN4 transmembrane proteins was observed in FB<sub>1</sub>-exposed Caco-2 cells showing the ability of FB<sub>1</sub> to impair the intestinal barrier integrity (Romero et al., 2016).

In our study no significant effect ( $P \ge 0.05$ ) was observed on TEER after 1, 2 or 24 h of Apical (Ap) or Basolateral (Bl) exposure to all the doses of FB<sub>1</sub> (0.5, 1.5, 3  $\mu$ M) (Fig. 4.3; 4.4). Our results are in agreement with previous studies (Wentzel et al., 2016) where Caco-2 cells were exposed for 24 h to different concentrations (0.5–45  $\mu$ M) of FB<sub>1</sub>. The results reported by Wentzel et al. (2016) showed that even at concentrations up to 45

 $\mu$ M, FB<sub>1</sub> did not affect cells. Similarly, exposure of Caco-2 cells to 0.001–10  $\mu$ M FB<sub>1</sub> revealed no cytotoxic effect of FB<sub>1</sub> at all the concentrations tested (Fernandez-Blanco et al., 2016).

Based on our findings FB<sub>1</sub> induced IL-8 release in Caco-2 cells even if only after Bl exposure to FB<sub>1</sub> at 3  $\mu$ M (Fig. 4.6), while no significant release of IL-6 or TNF- $\alpha$  was observed after Ap and Bl exposure to FB<sub>1</sub> at all doses (0.5, 1.5, 3  $\mu$ M) (Fig. 4.5).

The significant (P < 0.05) increase of IL-8 release induced by Bl exposure to FB<sub>1</sub> (3  $\mu$ M) (Fig. 4.6) was previously observed but after exposure to a higher concentration of FB<sub>1</sub> (17.2  $\mu$ M) (Minervini et al., 2014).

Regarding BEA its cytotoxicity was observed on Caco-2 cells at 20.6 and 12.7  $\mu$ M after 24 and 48 h exposure, respectively (Prosperini et al. 2012) and at 1 and 10  $\mu$ M (Fernandez-Blanco et al., 2016).

In our study a significant (P < 0.05) TEER decrease was observed after 1 h and 2 h of Bl exposure to BEA at 0.5 μM, whereas after 24 h of Bl exposure, BEA at 3 and 6 μM was found to significantly (P < 0.05) increase TEER (Fig. 4.9). When forced by an inducible promoter, an over-expression of OCLN leaded to a TEER increase of MDCK cells in a reversible manner, paradoxically increasing the paracellular flow of uncharged solutions (McCarthy et al., 1996). This paradoxical behavior still needs to be clarified (Barrett and Donowitz et al., 2001). Phosphorylation of TJ proteins, in particular CLDN, has also been shown to affect the epithelial barrier function (Stevenson et al. 1989; Findley et al., 2009). Moreover, protein kinase C (PKC) is involved in TJ regulation and has been demonstrated to act through an increase or decrease of TEER (Farhadi et al. 2006; Plotnikov et al., 2010).

The increase of TEER induced by BEA may be related to the modulation of TJ proteins, like claudin-2 and OCLN, with the effect of improving overall epithelial barrier function.

A significant (P < 0.05) release of IL-8 was induced by Ap exposure to BEA at 3 and 6  $\mu$ M and by Bl exposure to BEA only at 6  $\mu$ M (Fig. 4.11).

Concerning TNF- $\alpha$ , a significant release was observed after Ap exposure to 0.5 and 1.5  $\mu$ M BEA and after Bl exposure to 1.5  $\mu$ M BEA (Fig. 4.12).

In the present study the combined effects of FB<sub>1</sub> and BEA on Caco-2 cells were also evaluated.

After 24 h of Bl exposure, BEA alone at 3  $\mu$ M increased TEER (P < 0.05), but this effect was no longer observed when combined with FB<sub>1</sub> (Fig. 4.14), which shows no effect at 1.5  $\mu$ M after 1, 2 and 24 h of Ap or Bl exposure (P  $\geq$  0.05) (Fig. 4.13). These results suggest a possible interaction between these mycotoxins.

As previously discussed BEA alone was found to increase IL-6 and IL-8 release (Fig. 4.10; 4.11), but these increased releases were not observed when BEA was combined

with FB<sub>1</sub> (Fig. 4.15; 4.16), which induces IL-8 release at 3  $\mu$ M. Ap (0.5 -1.5  $\mu$ M) and Bl (1.5  $\mu$ M) exposure to BEA increased TNF- $\alpha$  release (Fig. 4.12), but not in the presence of FB<sub>1</sub> (Fig. 4.17).

TNF- $\alpha$  seems to play a key role in intestinal inflammation exerting a protective effect on the intestinal barrier function (Corridoni et al., 2012; Naito et al., 2003; Noti et al., 2010). The mechanisms of FB<sub>1</sub> toxicity include inhibition of ceramide synthase and accumulation of sphingoid bases, repression of protein kinase C and its down-stream effector TNF- $\alpha$  and mobilization of cellular calcium (Kim et al., 2006; Gopee et al., 2003). On the other hand BEA increases intracellular calcium and activates the calcium-dependent endonucleases and causes subsequent DNA fragmentation (Jestoi, 2008).

Only few studies so far have investigated the *in vitro* interactions between FB<sub>1</sub> and BEA, however synergic effects of these mycotoxins have been already reported (Klaric et al., 2007), even if the exact mechanism of the interaction needs to be clarified.

In this study FB<sub>1</sub> did not exert significant adverse effects on intestinal Caco-2 cells cultured on inserts, while BEA alone affected TEER and induced cytokine release but, in combination with FB<sub>1</sub>, these effects were no more observed. The disappearance of IL-6 and IL-8 release induced by BEA, may suggest a possible protective action of FB<sub>1</sub> (Cuzzocrea et al., 2008).

Our results suggest an interaction between FB<sub>1</sub> and BEA when combined on TEER and cytokine release of *in vitro* intestinal Caco-2 cells. Other studies are in progress to better understand these aspects.

## 4.5 References

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