

ORIGINAL ARTICLE

***In vitro* aflatoxin B₁ binding capacity by two *Enterococcus faecium* strains isolated from healthy dog faeces**

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Keywordsaflatoxin B₁, binding ability, *Enterococcus faecium*, food safety, healthy dogs.**Correspondence**

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2014/0682: received 31 March 2014, revised 18 October 2014 and accepted 10 December 2014

doi:10.1111/jam.12726

Abstract**Aim:** This study evaluated the binding capacity of aflatoxin B₁ (AFB₁) by two *Enterococcus faecium* strains (MF4 and GJ40) isolated from faeces from healthy dogs.**Materials and Methods:** The binding assay was performed using 50 and 100 ppb of AFB₁ analysing the effects of the viability, incubation time and pH on AFB₁ binding. Binding stability was determined by washing three times the bacteria-AFB₁ complexes with phosphate buffer saline.**Results:** Both GJ40 and MF4 strains have the ability to remove AFB₁ from aqueous solution. Viable cells were slightly more effective in AFB₁ binding than nonviable ones for both strains. *Enterococcus faecium* GJ40 removes 24–27% and 17–24%, and *Ent. faecium* MF4 removes 36–42% and 27–32% of AFB₁ (50 and 100 ppb, respectively) throughout a 48 h incubation period. In general, the removal of AFB₁ was highest at pH 7.00 for both strains. The stability of the bacteria-AFB₁ complex formed was found to be high (up to 50% of AFB₁ remained bounded in bacterial cell after three washes with phosphate buffered saline).**Conclusion:** The *Ent. faecium* strains assayed are capable of removing AFB₁ under different conditions *in vitro*.**Significance and Impact of the Study:** This is the first AFB₁ binding assay performed with *Ent. faecium* strains isolated from dog faeces, being an interesting strategy for AFB₁ decontamination of pet food.**Introduction**

Mycotoxins are food contaminants with harmful impact on human and animal health (CAST, 2003). Domestic dogs play several important roles in modern human society (Suchodolski *et al.* 2004). Pet food elaboration requires the use of appropriate technology to ensure its nutritional value and enhance life quality of pets. Many owners of pet food industries are concerned about the potential risk for mycotoxins contamination in pet foods, as the pets are fed for longer periods of time. Thus, these animals may become more vulnerable to chronic exposure to toxicants, especially to aflatoxins (AFs) which cause acute hepatotoxic and carcinogenic effects in dogs (Binder 2006; Boermans and Leung 2007). Previous works report important percentages of *Aspergillus* section *Flavi* AFs producers strains and samples contaminated

with aflatoxin B₁ (AFB₁) in ready-to-eat pet foods (Basalan *et al.* 2004; Leung *et al.* 2006; Campos *et al.* 2008, 2009; Fernández Juri *et al.* 2009a,b). Recently, international legislation establishes AFB₁ limits in pet food of 20–300 ppb in USA and 10–50 ppb in Europe (Aquino and Corrêa 2011). Several procedures are used for mycotoxins decontamination such as physical, chemical and biological methods (Park 1993). Live micro-organisms can decontaminate either by attaching the mycotoxin to their cell wall components or by active internalization and accumulation. Dead cells of microbes can also adsorb mycotoxins. This phenomenon can be exploited in the creation of biofilters for fluid decontamination or probiotics (which have proven binding capacity) to bind and remove the mycotoxin from the intestine (Juodeikiene *et al.* 2012). Yeast and lactic acid bacteria (LAB) cells can bind different molecules on the cell wall surface

(Bolognani *et al.* 1997; Santos *et al.* 2006). The mycotoxins removal is mainly produced by adhesion to cell wall components rather than by covalent binding or metabolism, as the dead cells do not lose their binding ability (Celyk *et al.* 2003; Santin *et al.* 2003; Baptista *et al.* 2004). Several LAB strains of food or animal origin have been tested for their ability to bind AFs and other mycotoxins to their surface (El-Nezami *et al.* 2002; Fuchs *et al.* 2008). Researchers have reported that different bacterial strains are able to bind AFB₁ in significantly different amounts (Peltonen *et al.* 2001). Fuchs *et al.* (2008) showed that LAB strains can remove patulin and ochratoxin A from aqueous solution in different levels. *Enterococcus faecium* is a member of LAB mostly found in nature and has various applications in the processing of some fermented dairy products (Giraffa 2003). Enterococci are used as probiotics in commercial formulation specially destined for animal feeding being mostly produced in foreign countries, not in Argentina. These strains produce antimicrobial substances against different bacteria as well as some fungal strains (Simonetta *et al.* 1997). In a previous work, the inhibitory effect of *Ent. faecium* strains isolated from the normal microbiota of healthy dogs gut on fungal growth parameters, and AFB₁ production by aflatoxigenic strains on *in vitro* assays was determined (Fernández Juri *et al.* 2011). The results obtained in the assay are promising due to the high percentages of AFB₁ production inhibition by *Ent. faecium* strains, even though they were unable to reduce *Aspergillus* strains growth rate. Topcu *et al.* (2010) reported the AFB₁ and patulin detoxification capacity of two commercial probiotic strains (*Ent. faecium* M74 and EF031) from aqueous solution. Both strains assayed had the ability to remove high percentages of the toxins throughout the assays. The viability of the bacteria did not have any significant effect on the detoxification of AFB₁ and patulin. However, there is no information available respect to AFB₁ binding capacity of *Ent. faecium* isolated from healthy dog faeces. Therefore, the aim of this study was to study the AFB₁ binding by two *Ent. faecium* strains isolated from faeces of healthy dogs that inhibit AFB₁ production in high percentages on *in vitro* assays. The effect of cell viability, incubation time and pH in the binding process and the stability of the AFB₁-viable cell were investigated.

Materials and methods

Chemicals and media

Man-Rogosa (MRS) (Britania) was used for *Ent. faecium* strains growth. Aflatoxin B₁ was obtained from Sigma-Aldrich (St. Louis, MO) and phosphate buffer solution

(PBS) from Britania (Buenos Aires, Argentina). For the AFB₁ binding assay, solid AFB₁ was suspended in benzene–acetonitrile (97/3, v/v) to obtain an AFB₁ concentration of 1000 ppb, then benzene–acetonitrile was evaporated by nitrogen stream and AFB₁ was resuspended in methanol (stock solution). Two working solutions (50 and 100 ppb) were prepared in PBS at pH 7.00 using the stock solution of AFB₁ in methanol (El-Nezami *et al.* 1998; Haskard *et al.* 2001).

Bacterial strains, culture conditions and count

Cultures of *Ent. faecium* MF4 and GJ40 were isolated from faeces of healthy dogs. Strains selection was based on their capacity of AFB₁ reduction or production inhibition (Fernández Juri *et al.* 2013). *Enterococcus faecium* strains were cultivated in MRS broth (24 h, 37°C) for preparation of overnight cultures. Samples of 1 ml of cultivated bacterial suspensions were decimally diluted in sterile peptone water (0.1%, w/v). Colony forming units (CFU) in overnight cultures were determined by plate counting on MRS Agar. In this study, all incubations were carried out at 37°C, and all centrifugations were at 3000 g for 15 min (4°C).

Mycotoxin binding assay

Viability assay

Cultures of each strain were divided in two aliquots. Aliquot 1: a volume of the culture broth (1×10^8 bacteria) was transferred into microtubes and centrifuged (viable cells). Aliquot 2: a volume of culture broth (1×10^8 bacteria) was transferred into microtubes then autoclaved at 121°C for 20 min (nonviable cells) and centrifuged. Supernatants were removed, and bacterial pellets (viable and nonviable cells) were washed with deionized water and centrifuged again (Peltonen *et al.* 2001; Topcu and Bulat 2010). For each strain, bacterial pellets (viable and nonviable) were resuspended in 2 ml of AFB₁ working solutions (50 and 100 ppb, pH 7.00), and the AF determination was performed after incubation at 24 h. Mycotoxin solutions without cells were used as controls. All the experiments were carried out by triplicate and repeated three times.

Incubation time effect

For incubation time effect, samples of *Ent. faecium* cells + AFB₁ were mixed in vortex shortly (5 s) and incubated for 48 h on an orbital shaker (New Brunswick Scientific CO., INC., Edison, NJ) with soft agitation. Supernatant samples (500 µl) were collected by centrifugation after incubation for 1, 24 and 48 h, and kept at –20°C for further analysis.

pH effect

For pH effect assay, PBS solution (pH 7.00) containing AFB₁ was adjusted to pH 3.00 and 5.00 with 1 mol l⁻¹ ClH and to 8.00 with 1 mol l⁻¹ NaOH. The effect of pH was tested after incubation at 25 °C for 24 h (El-Nezami *et al.* 1998).

Stability AFB₁-bacteria complex

The stability of the bacteria-AFB₁ complex was evaluated by determining the amount of AFB₁ remaining bound after three washes. The AFB₁ stability assay was performed at pH 3.00, 5.00, 7.00 and 8.00 after incubation at 24 h. Bacterial pellets with bound AFB₁ were suspended in 2 ml of PBS (pH 7.00) and incubated for 10 min at 25 °C. The bacteria were pelleted, and a volume of the supernatant was collected for the quantification of AFB₁ released from bacteria. This washing procedure was repeated another two times (Haskard *et al.* 2001; Peltonen *et al.* 2001).

Quantification of AFB₁ by HPLC

The quantification was performed by HPLC according to the methodology proposed by Trucksess *et al.* (1994) with some modifications (Cole and Dorner 1994), and the AFB₁ solutions (standards) were prepared according to AOAC (1995). An aliquot (200 µl) of the samples was derivatized with 700 µl trifluoroacetic acid: acetic acid: water (20 : 10 : 70, v/v). Chromatographic separations were performed on stainless steel, C18 reversed phase column (VARIAN, 150 × 4.6 mm id., 5 µm particle size). Water: methanol: acetonitrile (4 : 1 : 1, v/v) was used as mobile phase at a flow rate of 1.5 ml min⁻¹. The fluorescence of AFB₁ derivatives was recorded at excitation and emission wavelengths of 360 and 460 nm, respectively. The concentration of this toxin was quantified by correlating peak heights of sample extracts with those of standard curves. The detection limit of the analytical method was 0.1 ng g⁻¹. The percentage of mycotoxin bound to the bacteria was calculated using the equation: % Reduction = 100 × (1 - mycotoxin peak area of sample/mycotoxin peak area of control).

Statistical analysis

Data were analysed with an analysis of variance. Means were compared using a linear mixed model and Fisher's protected least significant difference (LSD) test to compare the AFB₁ binding along the treatments with AFB₁ in control assays. The analysis was conducted using software INFOSTAT, 2011 (Di Rienzo *et al.* 2011).

Results

Viability effect of *Enterococcus faecium* strains on AFB₁ binding

The viability of *Ent. faecium* cells after the preparation of two aliquots (aliquot 1: 1 × 10⁸ CFU ml⁻¹ of viable cells and aliquot 2: 10⁸ × 10 CFU ml⁻¹ of heat-killed cells) was determined by plate count method; 95% and 98% of *Ent. faecium* GJ40 and *Ent. faecium* MF4 cells in aliquot 1 remained viable, respectively. Both strains were 0% viable after aliquot 2 analyses (data not shown).

Figure 1 shows the effect of cell viability on AFB₁ binding capacity at pH 7.00 and 25 °C. Results showed no significant difference in AFB₁ removal between viable and nonviable cells in all treatments except for the interaction between *Ent. faecium* GJ40 at AFB₁ concentration of 50 ppb where the viable cells reduced the toxin concentration in a 20.55% and nonviable cell in a 15.45%, respect to the control ($P < 0.05$). The highest AFB₁ reduction was observed by *Ent. faecium* MF4 in the interaction assays using both toxin concentrations and both viable and nonviable cells; at 50 ppb viable cells reduced toxin concentration in 23.09% and nonviable cells in 25.51%. At 100 ppb, viable cells of *Ent. faecium* MF4 reduced AFB₁ in a 25.20% and nonviable cell in a 22.53%.

Incubation time effect on AFB₁ by *Enterococcus faecium* strains

Figure 2 shows the effect of incubation time on AFB₁ binding capacity by viable cells at pH 7.00 and 25 °C. In general, the removal of AFB₁ slightly increased with extended incubation time, independently of concentration, for both strains; otherwise, results showed that *Ent. faecium* MF4 removed AFB₁ along the experiment with more efficiency than *Ent. faecium* GJ40 ($P < 0.05$). In the binding assay using 50 ppb of AFB₁ as control, *Ent. faecium* MF4 showed the highest reduction percentage (42.23%) at 48 h, while under the same conditions *Ent. faecium* GJ40 reduced the toxin concentration in a 27.35%. There were no statistical differences in the reduction of AFB₁ at 50 ppb by *Ent. faecium* GJ40 along the incubation time ($P < 0.05$). At 100 ppb, both strains reduced significantly AFB₁ respect to the control ($P < 0.05$), but in different percentage, being MF4 strain the most efficient (31.64% at 48 h).

pH effect on AFB₁ binding by *Enterococcus faecium* strains

The effect of pH on AFB₁ reduction by viable bacterial cells of *Ent. faecium* strains is shown in Table 1. Toxin

Figure 1 Aflatoxin B₁ detoxification capacity of *Enterococcus faecium* strains at pH 7.00 and 25°C. (◐) % AFB₁ reduction by viable cells; (◑) % AFB₁ reduction by nonviable cells. SD: standard deviation. Values with the same letter are not significantly different according to least significant difference (LSD) test ($P < 0.05$). Statistical analysis compared means of the AFB₁ concentration of the phosphate buffered saline (PBS) as control and PBS with the toxin.

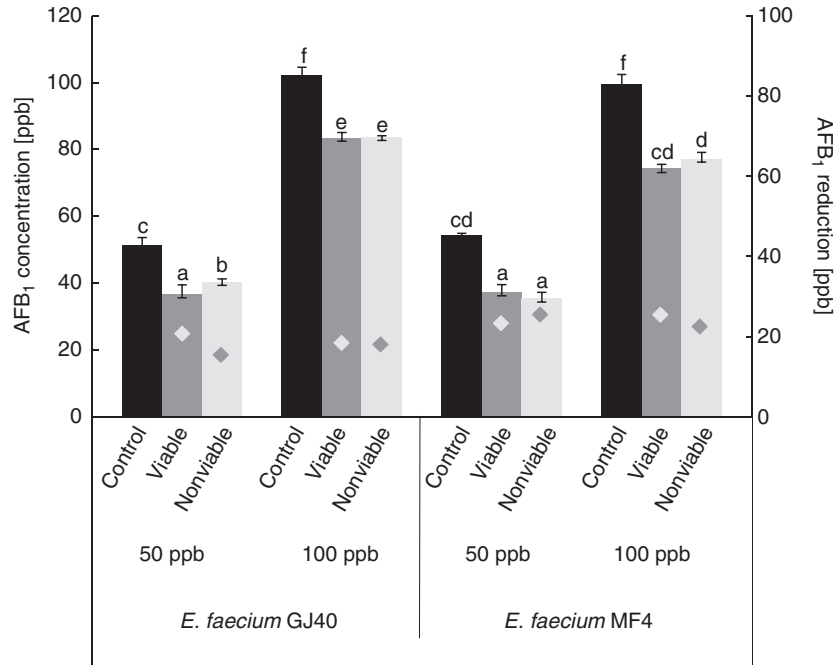
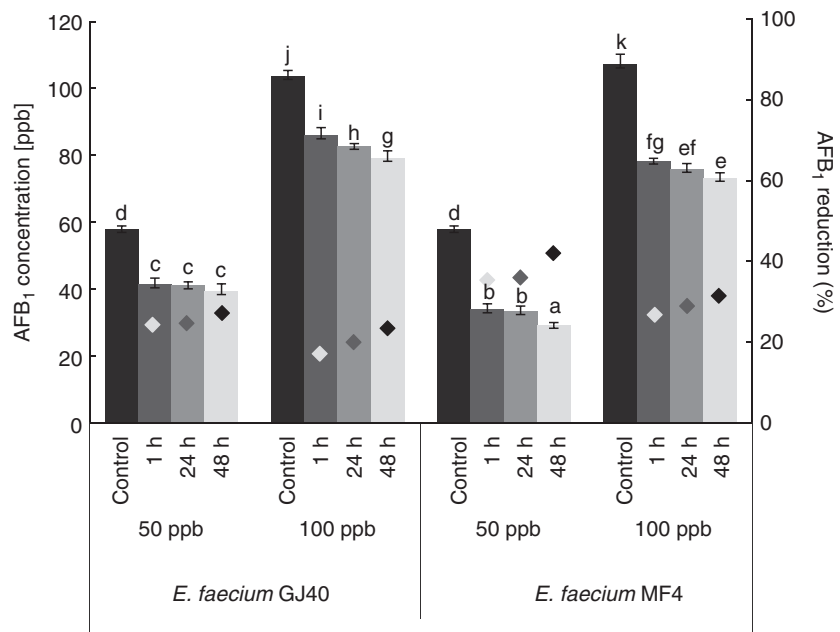


Figure 2 Effect of incubation time on AFB₁ detoxification by *Enterococcus faecium* viable cells at pH 7.00 and 25°C. (◐) % AFB₁ reduction at 5 h; (◑) % AFB₁ reduction at 24 h; (◒) % AFB₁ reduction at 48 h. Values with the same letter are not significantly different according to least significant difference (LSD) test ($P < 0.05$). Statistical analysis compared means of the AFB₁ concentration of the phosphate buffered saline (PBS) as control and PBS with the toxin.



reduction by viable bacterial cells varied in the range of pH analysed for both AFB₁ concentrations used in the assay. The highest reduction percentage was obtained by *Ent. faecium* MF4 at pH 7.00 in the assay using 50 ppb (28.78%) and the lowest by *Ent. faecium* GJ40 at pH 8.00 using 100 ppb (10.75%). In general, AFB₁ removal was major at pH 7 for both concentrations assayed by both *Ent. faecium* strains. No significant differences in AFB₁

binding capacity by *Ent. faecium* strains were detected. Significant differences were found between the concentrations assayed ($P < 0.05$).

Binding stability of bacteria-AFB₁ complex

Figure 3 shows the stability of the bacteria-AFB₁ complex after three washes with PBS at pH 7.00. Aflatoxin B₁ still

Table 1 Effect of pH on AFB₁ detoxification by *Enterococcus faecium* strains isolated from healthy dog faeces at 25°C for 24 h

	<i>Ent. faecium</i> GJ40				<i>Ent. faecium</i> MF4			
	50 ppb		100 ppb		50 ppb		100 ppb	
	(AFB ₁) [*] ± SD	AFB ₁ (%) [†]	(AFB ₁) [*] ± SD	AFB ₁ (%) [†]	(AFB ₁) [*] ± SD	AFB ₁ (%) [†]	(AFB ₁) [*] ± SD	AFB ₁ (%) [†]
PBS*								
pH 3.00 + AFB ₁	54.06 ± 2.00 ^{efg}	18.49^e	101.9 ± 2.2 ^{lm}	14.46^c	52.23 ± 3.48 ^{def}	26.80^j	101.13 ± 1.20 ^{lm}	20.56^f
pH 3.00 + AFB ₁ + bacteria	44.23 ± 1.05 ^{bc}		87.16 ± 1.75 ^{ij}		38.23 ± 0.45 ^{ab}		80.33 ± 0.51 ^{hlm}	
pH 5.00 + AFB ₁	57.06 ± 1.93 ^g	15.65^{cd}	104.86 ± 4.47 ^m	10.90^a	58.13 ± 1.95 ^g	24.47ⁱ	103.16 ± 2.15 ^{lm}	12.69^b
pH 5.00 + AFB ₁ + bacteria	48.13 ± 0.80 ^{cd}		93.43 ± 4.3 ^k		43.90 ± 1.25 ^{ab}		90.06 ± 0.80 ^{jk}	
pH 7.00 + AFB ₁	52.46 ± 2.45 ^{ef}	24.77ⁱ	101.62 ± 2.45 ^{lm}	20.22^f	56.13 ± 2.45 ^{fg}	28.78^k	101.6 ± 1.74 ^{lm}	21.06^g
pH 7.00 + AFB ₁ + bacteria	39.46 ± 0.83 ^a		81.06 ± 0.83 ^h		40.00 ± 0.83 ^{ab}		80.20 ± 1.05 ^h	
8.00 + AFB ₁	57.96 ± 2.00 ^g	10.75^a	99.63 ± 8.71 ^l	16.66^d	56.76 ± 3.97 ^g	22.25^{gh}	102.96 ± 3.70 ^{lm}	22.79^h
pH 8.00 + AFB ₁ + bacteria	51.73 ± 1.00 ^{de}		83.03 ± 2.02 ^{hi}		44.13 ± 1.42 ^{bc}		79.50 ± 0.81 ^h	

PBS, phosphate buffer solution adjusted at different pH values; SD, standard deviation.

Mean values based on triplicated data. Values with the same letter are not significantly different according to least significant difference test ($P < 0.05$).

Statistical analysis was performed separately for concentration and percentage values and statistical data must be read separately.

*Concentration of AFB₁.

†Percentage of AFB₁ reduction.

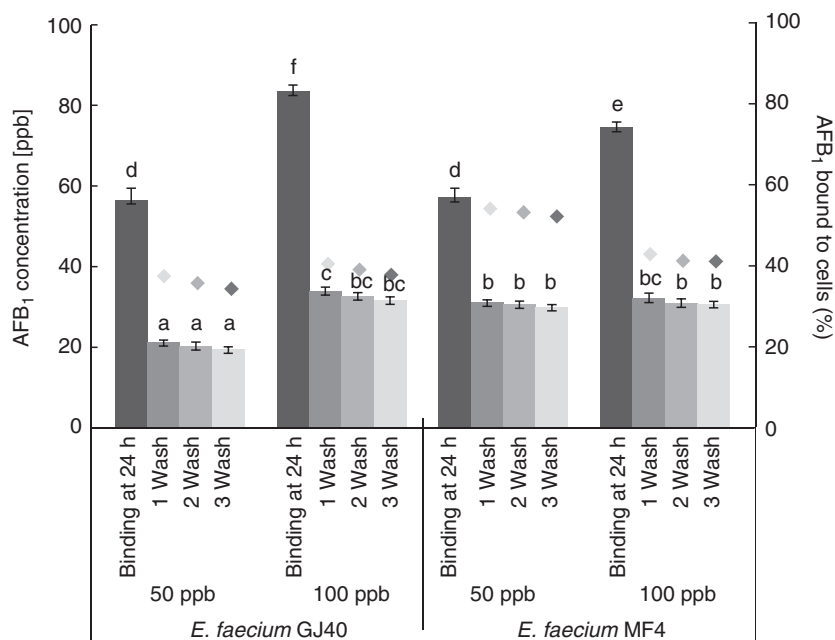


Figure 3 Stability of the complex viable bacteria-AFB₁ after three washes with phosphate buffered saline (PBS) at pH 7.00 after 24 h. (◐) % AFB₁ bound at one wash; (◑) % AFB₁ bound at two wash; (◒) % AFB₁ bound at three wash. Values with the same letter are not significantly different according to least significant difference (LSD) test ($P < 0.05$). Statistical analysis compared means of the AFB₁ concentration in the binding at 24 h and AFB₁ concentration remained in the cell after the washing procedure.

remains bound in the cell of *Ent. faecium* GJ40 in a 34–37% and in 41.31–52.62% in *Ent. faecium* MF4 after three washings with PBS. In all cases, no significant differences in the release of the toxin were found after the first wash ($P < 0.05$). The pH of washing solution (3.00, 5.00 and 8.00) did not affect the binding stability of bacteria-AFB₁ complex (data not shown).

The LSD test of data shows the significant differences among control treatments and interacting assays. The statistical analysis of strains showed that all analysed

factors influenced significantly on AFB₁ binding ($P < 0.05$) (Table 2).

Discussion

The present study shows the capacity of removing 50 and 100 ppb of AFB₁ in aqueous solution by two *Ent. faecium* strains isolated from healthy dog faeces. Viable and non-viable cells of both strains (GJ40 and MF4) removed AFB₁ in different percentages along the experiments, but

Table 2 Analysis of variance of effect of strains (S), AFB₁ concentration (C), viability (V), incubation time (I) pH (P) and their interactions on AFB₁ binding

Source of variation	df	AFB ₁ binding	
		MS	F
S	1	94.74	23.99*
C	1	5319.27	1347.03*
V	2	1342.54	339.98*
I	3	1751.59	610.98*
P	3	158.54	23.52*
S × C × V	2	11.60	2.94*
S × C × I	3	3.00	1.05*
S × C × P	3	2.41	0.36*

df, degrees of freedom; MS, mean square, F, F-Snedecor.

*Significant $P < 0.05$.

Ent. faecium MF4 was more effective in the removal of the toxin. The results obtained are similar of those obtained by Topcu *et al.* (2010), who assayed the binding capacity of two probiotic *Ent. faecium* strains. The researchers found that the strains were able to bind AFB₁ in high percentages under different conditions of pH, viability and incubation time (46%).

In the viability assay, no significant differences were found in AFB₁ binding by viable and nonviable *Ent. faecium* GJ40, but the toxin removal by *Ent. faecium* MF4 viable was significantly different respect to the nonviable cells, even though the reduction percentage was similar. According to this, it can be considered that detoxification of AFB₁ by *Ent. faecium* may be due to the binding of the mycotoxin to the bacterial cell wall components. This mechanism has been studied and also postulated by other authors (El-Nezami *et al.* 1998; Haskard *et al.* 2000, 2001). The cell wall peptidoglycans and polysaccharides have been proposed to be responsible for mycotoxin binding by bacteria (Haskard *et al.* 2001; Lahtinen *et al.* 2004). In several previous studies, it has been reported that heat treated bacteria were more effective in removing AFB₁ than viable cells (Shetty and Jespersen 2006; Topcu and Bulat 2010); the results obtained in the present work are similar to another studies reporting that no significant differences were found in the removal of the toxin by viable and nonviable cells (El-Nezami *et al.* 1998; Haskard *et al.* 2001; Pizzolito *et al.* 2011). Therefore, the fact that nonviable and viable micro-organisms are able to remove AFB₁ *in vitro* conditions with similar efficiency suggests that the removal process does not require metabolic conversion of the toxin by cells and it could be dependent on the strain. Additionally, these results indicate that the inclusion of viable or nonviable micro-organisms in the diet of animals would be equally effective against aflatoxicosis. This is very important because

the use of nonviable cells decreases the potential pathogenic risks of their inclusion in the diet. Besides, the use of viable cells is an interesting point as the assayed *Ent. faecium* strains have benefic properties for the host, such as antimicrobial activity (Fernández Juri *et al.* 2013).

About the effect of incubation time, Peltonen *et al.* (2001) reported that the removal of AFB₁ was a fast process and the AFB₁ binding by *Lactobacillus amylovorus* CSCC 5197 increased from 52 (0 h) to 68% (48 h). In another study, it was observed that at zero hour the percentage of removed AFB₁ was not significantly different from the one at 72 h (Pizzolito *et al.* 2011). On the other hand, Topcu *et al.* (2010) reported that at 1 h, *Ent. faecium* M74 and EF031 strains removed AFB₁ which was approximately 65% of the total of AFB₁ removed throughout the whole incubation period of 48 h. In the present work, both strains at both concentrations detoxified AFB₁ quickly at 1 h, but the removal maintained similar up to 48 h of assay, even though there were significant differences. These results suggest that the binding of the toxin occurs quickly, but does not increase the removal considerably with the incubation time.

Regarding to pH effect, the highest reduction percentage was obtained by *Ent. faecium* GJ40 at pH 7.00. Respect to *Ent. faecium* MF4, pH values assayed (3.00, 5.00, 7.00 and 8.00) did not affect statistically the removal of the toxin ($P < 0.05$), except at pH 5 using 100 ppb. These results partially agree with those previously reported. Some researchers reported that binding process is not exclusively dependent of pH. Zinedine *et al.* (2005) reported that all the *Lactobacillus* spp. assayed removed the toxin from 5% to 40% when pH increased from 3 to 5.5. On the other hand, Rayes (2013) reported that the highest reduction percentage of AFB₁ by a pool of probiotic LAB occurred pH 8.5 while the lowest removal was at pH 4.5. When the pool was combined with a *Saccharomyces cerevisiae* strain, the highest removal was at pH 4.5 and the lowest removal was at pH 8.5. In contrast, results in the binding capacity assay using *Lactobacillus rhamnosus* strains at pH 3 and 6 showed that there were no significant differences in the toxin reduction between any bacterial cell conditions (in solution, spray or freeze-dried) (Bovo *et al.* 2014). Similarly, Peltonen *et al.* (2001) assayed 12 *Lactobacillus*, five *Bifidobacterium* and three *Lactococcus* strains and concluded that the differences in AFB₁ binding by the strains were probably due to different bacterial cell wall and cell envelope structures. So, it can be said that the pH and incubation time dependence of AFB₁ binding vary between bacterial strains and their cell wall components.

Respecting the toxin concentration effect, Rahayu *et al.* (2007) demonstrated that increasing AFB₁ concentration

in aqueous medium did not affect the percentage of AFB₁ binding; however, it affects the binding speed. In addition, Pizzolito *et al.* (2011), in a binding assay with yeast and LAB using different AFB₁ concentrations, reported that the removal of the toxin depended of the strain; some were more effective at a lower concentration (*Lact. rhamnosus* I at 50 ppb) and some others were more effective at a higher concentration (*Lactobacillus acidophilus* 24 at 100 ppb and *Lactobacillus casei* subsp. *rhamnosus* at 500 ppb). The results obtained in the present work are similar to the mentioned above, as *Ent. faecium* strains were able to bind AFB₁ in higher percentages in the assay using 50 ppb of AFB₁, but this ability depends on the strain.

Related to stability of the viable cell-AFB₁ complex, results suggest that binding is a fast and reversible process. On the other hand, it could be proposed that metabolic conversion of the toxin by cells did not take place, because AFB₁ was released in the same chemical form from bacterial cells. Several researchers have previously reported the reversibility of the process; Haskard *et al.* (2001) analysed 12 LAB strains in both viable and nonviable forms, exhibiting reversible binding of AFB₁ after five washing steps with water (6–11%). Hernandez-Mendoza *et al.* (2009) reported that using PBS solution, about 60–70% of AFB₁ remained bound to the cells, suggesting that the toxin is attached to the bacteria by weak, no covalent interactions that could be at least partially reversible. Pizzolito *et al.* (2011) reported that after five washings with PBS, AFB₁ bound to different LAB cells were close to 50%, and the washing time did not change the release percentages when it varied from 1 to 60 min. Similarly, the results obtained in the present work showed that the toxin remained in the bacterial cells in a high percentage (35–50%) after three washes with PBS solution. According to this, it could be suggested that neither the entrance of AFB₁ into the cell nor its metabolic conversion is necessary; therefore, micro-organism cell wall components may be involved in AFB₁ removal, as was suggested by several authors (Lahtinen *et al.* 2004; Karaman *et al.* 2005). These results may be attributed to differences of cell wall structure characteristics of each strain.

Concluding, the results obtained in the present work show that *Ent. faecium* GJ40 and *Ent. faecium* MF4 assayed possess the capacity of removing AFB₁ from aqueous solutions with different efficiency under the assayed conditions. Even though the reduction percentages of AFB₁ in general were below 30%, these strains could be used as additives in dry pet food naturally contaminated with low levels of this toxin (Campos *et al.* 2009; Fernández Juri *et al.* 2009b). In addition, these strains also present benefic properties previously assayed

(Fernández Juri *et al.* 2013). Further experiments simulating the intestinal conditions are needed to determine the efficiency of toxin removal and once the bacteria are added to the pet food.

Acknowledgements

This work was carried out with grants from Secretaría de Ciencia y Técnica Universidad Nacional de Río Cuarto (SECYT-UNRC), Consejo Nacional de Ciencia y Técnica (CONICET) and Fondo para la Investigación Científica y Tecnológica (FONCYT-PICT) (Argentina). M.G. Fernández Juri, A.M. Dalcerro and C.E. Magnoli are members of the research career of CONICET.

Conflict of Interest

The authors of this manuscript declare that they have no conflict of interest.

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