



Biological reduction of aflatoxin B1 in yogurt by probiotic strains of *Lactobacillus acidophilus* and *Lactobacillus rhamnosus*

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Received: 27 May 2019 / Revised: 26 November 2019 / Accepted: 3 December 2019 / Published online: 23 December 2019
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Abstract The present study was conducted to investigate the ability of two probiotic strains, *L. acidophilus* PTCC 1643 and *L. rhamnosus* PTCC 1637, to bind aflatoxin B₁ (AFB₁, 20 ng/ml) in comparison with yogurt starter cultures, at equal bacterial count ($\sim 10^9$ LogCFU/ml) during a 21-day storage period at 4 °C. All assessed treatments exhibited high percentages of AFB₁-binding, ranged from 64.56 to 96.58%. However, the ability of probiotic bacteria was statistically higher than yogurt starter cultures. Aflatoxin binding ability of the selected lactic acid bacteria was dependent on both time and bacteria species. The highest and the lowest percentages of AFB₁ removal was observed at 11th day of cold storage by *L. rhamnosus* ($96.58 \pm 3.97\%$) and at the first day of storage for yogurt starter cultures ($64.56 \pm 5.32\%$), respectively. The stability of bacterial cells-AFB₁ complex was remarkable, since only 0.84–26.75% of bounded AFB₁ was released from bacterial cells after 3 times washing during the storage period.

Keywords Aflatoxin removal · Probiotic yogurt · Binding ability · Complex stability

Introduction

The human diet may contain a miscellaneous array of natural mutagenic or carcinogenic compounds, due to the pollution of raw materials or the formation of toxic metabolites throughout food processing, cooking or storage (Osowski et al., 2010; Wang et al., 2011). Generally, mycotoxins are naturally occurring poisonous secondary metabolites of filamentous fungi and mainly produced by *Aspergillus*, *Penicillium*, and *Fusarium* species (Priyanka et al., 2014; Venkataramana et al., 2014). These toxic compounds play an indubitable performance in the reduction of the marketable and hygienic quality of various products (Dalié et al., 2010). The FAO (Food and Agriculture Organization of the United Nations) assessed that more than 25% of the world agricultural production is polluted by mycotoxins (Marin et al., 2013). Mycotoxin contamination of the food chain has a drastic impact and uncountable economic costs. However, the United States Food and Drug Administration (FDA) based on computer modeling, evaluated that the potential economic costs of crop losses due to aflatoxins, fumonisins and trichothecenes in the USA are expected to be \$932 million per year (Milićević et al., 2010). In the same way, the large world consumption of the most important foodstuffs, such as corn (1033.7 million tonnes/year), milk (876 million tonnes/year), wheat (757.9 million tonnes/year), soybean (336.7 million tonnes/year) and peanut (45.45 million tonnes/year), makes the presence of mycotoxins as a serious problem in these products.

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Aflatoxins (AFs), are highly toxic mycotoxins produced by some *Aspergillus* species especially *A. flavus*, *A. parasiticus* and rarely *A. nomius*. Currently, there are 20 related compounds defined by the term of AF (Prandini et al., 2009). Among them, AFB₁ and AFB₂ are produced by *A. flavus*, but AFG₁ and AFG₂ along with AFB₁ and AFB₂ are produced by *A. parasiticus* (Bennett and Klich, 2003; Kumar et al., 2017). Besides, AFM₁ and AFM₂ are the hydroxylated metabolites of AFB₁ and AFB₂, respectively in lactating animals and humans (Hussain and Anwar, 2008). Intake of aflatoxin contaminated foods and feeds could lead to acute and chronic aflatoxicosis, including carcinogenic, mutagenic, teratogenic, neurotoxic, oestrogenic, and immune suppressive effects (Groopman et al., 2008; Ishikawa et al., 2017; Jiang et al., 2015; Milićević et al., 2010; Sellamani et al., 2016; Smith et al., 2017; Sun et al., 2018). International Agency for Research on Cancer (IARC) classified AFB₁ as a group I, carcinogen for humans (IARC, 2016).

Due to the harmful aftermaths of mycotoxins, certain procedures have been established to inhibit the development of these compounds and as well, to detoxify human foodstuffs and animal feedstuffs (Hathout and Aly, 2014; Kabak et al., 2006). These methods include: (1) the inhibition of mycotoxin pollution, (2) the decontamination/detoxification of foods and feedstuffs contaminated with mycotoxins, and (3) prevention of absorption of mycotoxin content of consumed food into the digestive tract (Hathout and Aly, 2014). Several methods, including chemical, physical and biological control strategies have been proposed and implemented to decrease level of aflatoxins in foods and feeds with varying degrees of successes (Abdallah et al., 2015; Bozoğlu, 2009; Karlovsky et al., 2016; Shao et al., 2016; Zaki et al., 2012). Among these methods, chemical and physical procedures have some limitations, such as concerning safety issues, losses in the nutritional value, altered organoleptic characteristics of the products, limited efficacy and cost implication (Gowda et al., 2007; Guan et al., 2011; Méndez-Albores et al., 2005; Puzyr et al., 2010). Therefore, it is crucial to attain innovative toxin removal or detoxifying approaches, especially for elimination of aflatoxins, to promote food safety.

Probiotics are described as “living microorganisms which when ingested in adequate amounts, beneficially influence the health of the host by improving the composition of intestinal microbiota” (FAO/WHO, 2002). Probiotics have shown physiological function and promote the body’s immunity (Nooshkam et al., 2018; Shah, 2000a; Yerlikaya, 2014). Yogurt, the best carrier of probiotics, traditionally is produced using *Streptococcus thermophilus* and *Lactobacillus delbrueckii* ssp. *bulgaricus* as starter cultures. Nowadays, an adequate level of lactic acid bacteria (LAB), particularly probiotic strains are considered as

a key elements of yogurt quality (Ziarno and Zaręba, 2019). Several researchers have tested the lactic acid bacteria, as *Lactobacillus* and *Bifidobacterium* species for their ability to bind AFB₁ both in vitro and in vivo conditions, due to their GRAS (generally recognized as safe) status and use as probiotics (Liew et al., 2018; Zoghi et al., 2014). For example, Peltonen et al. (2001) revealed that a properly high efficiency (more than 50%) of binding of aflatoxin B₁ from a buffered saline solution is displayed by strains of *Lactobacillus amylovorus* and *Lactobacillus rhamnosus*, while *Lactococcus lactis* spp. *cremoris*, *Bifidobacterium animalis* and *Bifidobacterium lactis* were slightly less effective (40–50%). Corassin et al. (2013) compared the AFB₁ binding capacity of *L. delbrueckii* ssp. *bulgaricus*, *L. rhamnosus*, and *B. lactis* in combination with heat-inactivated *S. cerevisiae*. This assimilation certified complete mycotoxin binding (100%).

The hot and humid climates and storage condition of many food products as milk powders, polluted and unhygienic environment, frequent opening of bags contain these products, using unclean cups or containers for measurement, as well as, unclean and unsterilized packaging materials may lead to the growth of *Aspergillus* species, particularly *Aspergillus flavus* and subsequently aflatoxins production. Among aflatoxins, AFB₁ is the most toxic metabolite as compared to other mycotoxins. Therefore, the main objectives of this investigation were to: (1) determine the AFB₁-binding ability of the selected probiotic bacteria in comparison with yogurt starter cultures and (2) to evaluate the stability of the complexes (bacterial cells-AFB₁) during 21-days cold storage period.

Materials and methods

Chemicals and media

Aflatoxin B₁ (AFB₁, from *Aspergillus flavus*) in powder form was purchased from Sigma-Aldrich, Germany. All the solvents used include methanol, acetonitrile, benzen and n-hexane were obtained from Merck (Germany) and dichloromethane was purchased from Samchun (Korea). For bacterial cultivation and counting, de Man–Rogosa–Sharpe (MRS) broth and agar (Liofilchem, Italia), peptone water (Merck) and bile salt powder (Merck) were used. Commercial Direct Vat Set (DVS) lyophilized pouches of yogurt starter cultures contain a mixed culture of *Streptococcus thermophilus* and *Lactobacillus bulgaricus* were supplied by Chr. Hansen company (Denmark). The lyophilized cultures were maintained according to the manufacturer’s instructions, at – 20 °C. The probiotic lactic acid bacteria strains, *Lactobacillus acidophilus* PTCC 1643 and *Lactobacillus rhamnosus* PTCC 1637, as lyophilized

ampoules were procured from Iranian Research Organization for Science and Technology (Persian Type Culture Collection, PTCC), Tehran, Iran.

Preparation of bacterial strains and growth conditions

L. acidophilus and *L. rhamnosus* were selected based on their use in various foodstuff, especially dairy products, and on their ability to bind different kinds of food contaminant, such as mycotoxins and heavy metals (Bhakta et al., 2012; Haskard et al., 2001; Zoghi et al., 2014). *L. rhamnosus* was inoculated directly into MRS broth, and *L. acidophilus* activated in MRS broth supplemented with 0.05% cysteine hydrochloride, then incubated (BINDER GmbH, Model KB 23, Germany) without shaking at 37 °C for 48 h, in aerobic and anaerobic conditions, respectively. The evaluation of bacterial cells concentration in the cultures were determined by conventional agar plating technique using MRS agar and further incubation of cultured plates at 37 °C for at least 48 h. At the end of the incubation period, bacterial counts were expressed as colony forming unit (CFU) per milliliter of the media. In order to storage, bacterial cells were kept at - 80 °C in MRS broth having 20% (v/v) glycerol, as cryoprotectant. At the time of experiments, strains were recovered from media, grown in MRS broth again and incubated, as previously described. After incubation, cells were collected by centrifugation at 6000 g for 10 min at 10 °C. Finally, bacterial pellets removed from the supernatant under the sterile conditions and washed once by sterile deionized water prior to use (El-Nezami et al., 1998; Elsanhoty et al., 2014).

Preparation of AFB₁ standard solution

AFB₁ powder was suspended in a mixture of HPLC grade acetonitrile/benzene (3:97 v/v) to obtain the approximate concentration of 10 µg/ml. Then, the standard solution was achieved by diluting the mixture in phosphate buffer saline (PBS) solution. The solvents were evaporated using a water bath (Memmert, Model WNB 14, Germany) by heating at 80 °C for 10 min (Haskard et al., 2001). The concentration of the standard solution was finalized and computed by the Lambert-Beer equation ($A = \epsilon cl$) using the absorbance (A) at 354 nm, a molar absorptivity $\epsilon_{354} = 19.950$ per Mol.cm and the optical distance crossed by light in the medium (l) (Zinedine et al., 2005). The obtained solution was transferred to a dark glass bottle and kept in refrigerator until used. An aliquot of this standard solution was diluted in PBS (pH ~ 7) to final concentration of 20 ng/ml and to perform the AFB₁ binding assay. For drawing of calibration curve, standard solutions of AFB₁ with different concentrations were prepared by dilution in acetonitrile,

water and methanol (20:50:30 v/v/v, respectively). Calibrations curve were obtained by plotting the peak area for each calibration solution against the concentration of AFB₁ injected (Sarлак et al., 2017). The correlation coefficient R^2 was 0.9997.

Preparation of yogurt

Set-type yogurt samples were made according to the method of performed by Elsanhoty et al. (2014) with some modifications. Reconstituted milk was prepared by diluting 10 g of skimmed milk powder into 100 ml sterile deionized water. The milk was then stirred for 5 min and heated at 90 °C for 15 min to both destroy pathogen microorganisms and raise the denaturation of whey proteins. Next, the milk cooled down to incubation temperature (45 °C) and inoculum was added. As the minimum concentration of 10⁸ CFU/ml of LAB is required for adequate and rapid binding of aflatoxins to the bacterial cell wall (Kabak and Var, 2008), the LAB strains was added so that the number of bacteria reaches to ~10⁹ CFU/g. Different batches from yogurt were produced as: negative control (NCON), milk without AFB₁ and probiotic strain and only inoculated with 0.05% yogurt starter cultures (~10⁹ CFU/g); positive control (PCON), milk with 20 ng/ml AFB₁ and without probiotic strain inoculated with 0.05% yogurt starter cultures; *L. acidophilus* probiotic yogurt (LAPY), milk with 20 ng/ml AFB₁ inoculated with 0.025% yogurt starter cultures (~5 × 10⁸ CFU/g) and *L. acidophilus* probiotic strain (~5 × 10⁸ CFU/g); *L. rhamnosus* probiotic yogurt (LRPY), milk with 20 ng/ml AFB₁ inoculated with 0.025% yogurt starter cultures and *L. rhamnosus* probiotic strain (~5 × 10⁸ CFU/g). Therefore, the initial viable cell counts of all the yogurts (probiotics and non-probiotics) were similar (~10⁹ CFU/g). All batches were incubated at 42 °C until reaching pH approximately to 4.60, at which the fermentation was terminated. The fermentation time to reach pH 4.60 for yogurt samples was about 4 h. Then, samples were immediately cooled and stored at 4 °C for 21 days. All analyses of the yogurt samples were performed in triplicate after production, and 1, 11- and 21-days during storage at refrigeration temperature.

Evaluation of pH and titratable acidity

The pH of the samples was measured using a digital pH meter (Metrohm Company, model 827, Switzerland) by direct immersion of the electrode in samples at room temperature. Titratable acidity was assessed via blending 9 g of sample with 9 ml of distilled water and titrating with 0.1 N NaOH using phenolphthalein (1% (w/v) in ethanol) as an indicator, to an end point of stable faint pink color for

30 s (Jooyandeh et al., 2015). Titratable acidity was asserted as a percentage of lactic acid.

Evaluation of syneresis

Syneresis was obtained by slight modification of the technique stated by Akgun et al. (2018). To measure syneresis, 5 g of yogurt weighed in centrifuge tubes and then centrifuged (HERMLE Labrotechnik GmbH, Model Z 206 A, Germany) at 2500 rpm for 10 min at 25 °C. The collected liquid from the sample that separated in the top of tube was then gently poured off, weighed and regarded as syneresis.

$$\text{Syneresis (\%)} = \frac{\text{Volume of upper phase}}{\text{Sample weight}} \times 100$$

Microbiological analysis

After serial dilution, bacteria were counted by the surface culture method. MRS-bile agar medium was used for the selective enumeration of probiotic bacteria (Kabak and Var, 2008).

Preparation of samples for AFB₁ binding assay

For each sample, after gentle mixing, 5 g yogurt was transferred in falcon tubes. Subsequently, 40 ml of dichloromethane was added, and suspension was agitated for 15 min and filtered. Thereafter, 10 ml of the filtrate was heated at 60 °C and after evaporation, the residual oily was re-suspended in a mixture of 0.5 ml PBS, 0.5 ml methanol and 1 ml of n-hexane. The suspension was centrifuged (Eppendorf, Model AG 22331, Germany) at 3500 g for 15 min at 10 °C. After removing n-hexane (upper layer), 100 µl of the aliquot were diluted with 400 µl of deionized water and 100 µl of the diluted samples was experimented for the AFB₁ assessment (El Khoury et al., 2011; Elsanhoty et al., 2014).

AFB₁ binding assay by HPLC

The HPLC technique used for the analysis of unbound AFB₁ present in the supernatant was carried out according to Soares et al. (2010) with minor modifications. The HPLC system (KNAUER smartline, Germany) equipped with a programmable fluorescence detector (RF-10AXL) and a pump solvent delivery system (model 1000). Separation was achieved by using a Eurospher 100-5 C18 reversed phase column (4.6 × 150 mm, 5 µm particle size, KNAUER Smartline) at 40 °C with an injection volume of 100 µl. Water–methanol–acetonitrile (5:3:2 vol/vol/vol) was used as the mobile phase, with a flow rate of 1 ml/min.

The excitation and emission wavelengths were set at 365 and 460 nm, respectively. Under these conditions, the retention time of AFB₁ was around 12 min. The percentage of the bound AFB₁ by the examined strains suspension was determined using the following equation:

$$\text{AFB}_1 (\%) = 1 - \frac{\text{AFB}_1 \text{ peak area of sample}}{\text{AFB}_1 \text{ peak area of toxin control}} \times 100$$

Stability of the bacterial cells-AFB₁ complex

The amounts of AFB₁ released from the bacteria-AFB₁ complexes were investigated after repetitive washes (three times) by HPLC. For determination of bacterial cells-AFB₁ complex stability, 1 g yogurt sample was mixed with 1.5 ml PBS solution for 15 s. Afterward, the mixture was kept at 37 °C for 5 min. For removing the bacterial cell, the suspension was centrifuged (Eppendorf, Model AG 22331, Germany) at 2000 g for 10 min at 10 °C s. Amount of AFB₁ in the supernatant (at the third washing stage) was determined using HPLC (Peltonen et al., 2001; Utami et al., 2017).

Statistical analysis

Data were analyzed as a completely randomized factorial design. The significance of the difference between means was characterized by Duncan's multiple range test ($p < 0.05$) using SPSS statistical software (version 20, SPSS Inc., USA). All statistics were the averages of triplicate trials, and the values were shown as the mean values. All graphs were created using Microsoft Excel 2013 software (Version 6.2, Palisade Corporation, New York, USA).

Results and discussion

pH and titratable acidity

The changes in pH of the tested samples after completion of the yogurt fermentation and during the 21-day cold storage (4 °C) are shown in Fig. 1. The average pH values for all samples ranged from 4.07 to 4.67 during the storage period. The highest pH value (4.67 ± 0.05) was recorded for PCON just after yogurt production and the lowest pH value (4.07 ± 0.09) was recorded for LOPY sample at the end of storage. PCON demonstrated greater pH value (4.54 ± 0.07) than other samples at the all storage periods. However, there was no significant difference ($p > 0.05$) between the pH values of non-probiotic yogurts, i.e. PCON and NCON samples. Furthermore, probiotic yogurts had noticeably lower pH values than positive and negative controls ($p < 0.05$). Among probiotic yogurts, LOPY yogurts had generally lower pH value than LOPY

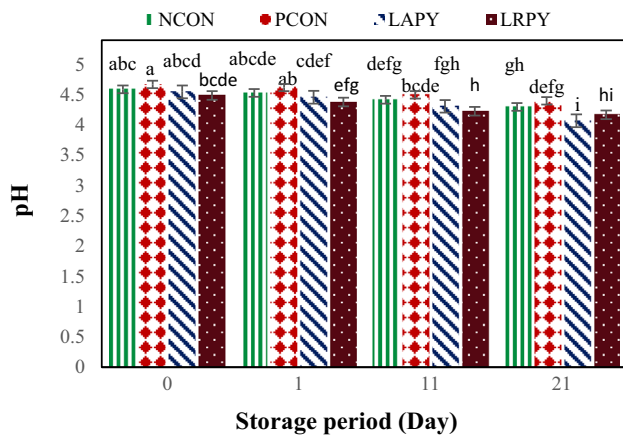


Fig. 1 Changes in the pH levels of yogurt samples during the cold storage. Different lowercase letters indicate significant differences at $p < 0.05$

samples but these differences were not significant ($p > 0.05$).

During storage period, the pH values of all yogurt samples declined significantly ($p < 0.05$). Nevertheless, pH values did not decrease lower than 4.0, which is usually considered unfavorable to the survival of probiotic bacteria (Yilmaz-Ersan and Kurdal, 2014). Beal et al. (1999) stated post acidification is the main reason of pH reduction during storage due to permanent metabolic activity of the starter cultures added to the product (mainly *L. delbrueckii* subsp. *bulgaricus*). This decrease in pH, is attributed to the use of residual carbohydrates by viable bacteria and production of several metabolites such as lactic acid, small amounts of CO_2 and formic acid from lactose (Panesar and Shinde, 2012).

The pH reduction was slightly more in probiotic yogurts in comparison with control samples. The pH varied between 4.07 ± 0.09 and 4.55 ± 0.11 for the probiotic batches of yogurt and 4.30 ± 0.03 and 4.67 ± 0.05 for the control batches during the cold storage. These results were similar to the findings of Shaghghi et al. (2013) who reported the lower pH for probiotic yogurts during the storage.

Like pH values, the titratable acidity (TA) values of yogurt samples during storage period were changed significantly (Fig. 2). The TA values of control and probiotic samples increased meaningfully ($p < 0.05$) throughout storage time and varied from 0.8 ± 0.07 to $1.77\% \pm 0.13$. The increase in TA values of fermented milks during cold storage is a common phenomenon. In general, the higher TA were obtained with probiotics yogurt samples, i.e. LOPY and LRPY samples in comparison with controls. These finding were in agreement with Yilmaz-Ersan and Kurdal (2014) who reported the higher level of TA for probiotic yogurts but were in contrast with Güler-Akin and

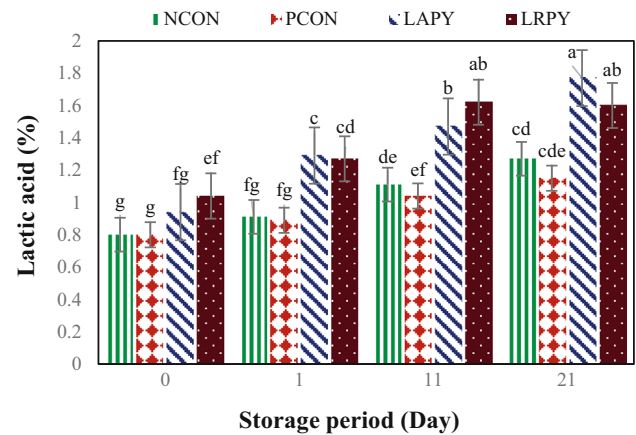


Fig. 2 Changes in the titratable acidity percentages of yogurt samples during cold storage. Different lowercase letters indicate significant differences at $p < 0.05$

Akin (2007) who described the lower level of TA in probiotic yogurts due to growth inhibition of *L. bulgaricus* by probiotic bacteria. Many factors could influence the acidity/pH values of yogurts during storage period which from those, the type of starter cultures, manufacturing methods and storage conditions are the more important. However, it should be considered that yogurt starter cultures can only produce lactic acid while *L. acidophilus* and *Bifidobacteria* have capability to produce both lactic and acetic acids (Yilmaz-Ersan and Kurdal, 2014). As it is shown in Figs. 1 and 2, PCON yogurt samples contained 20 ng/ml AFB_1 had slightly ($p > 0.05$) higher pH and lower TA in comparison with NCON. These findings indicate that addition of AFB_1 to yogurt may have negative effect on starter cultures activity.

Syneresis

Syneresis, is one of the main problems of yogurt industry. This is the expulsion of whey from three-dimensional casein networks, which turn out to be observable on the surface. Whey loss measures the level of collapsed gel and is an indicator for poor quality and stability. As it is shown in Fig. 3, control yogurt samples had lower syneresis in comparison with probiotic yogurts throughout the storage period. However, except at day 11, these differences were not significant ($p > 0.05$). The syneresis in the control samples during cold storage was in the range of 13.76 ± 0.12 – $26.57 \pm 0.09\%$, while in the probiotic yogurts it was varied in the range of 16.29 ± 0.10 – $27.22 \pm 0.11\%$. This may be due to the lower pH and the higher acidity in the probiotic yogurts in comparison with control samples, as a decrease in pH value in yogurt accelerates the syneresis (Athar et al., 2000). The highest syneresis was determined in LRPY sample

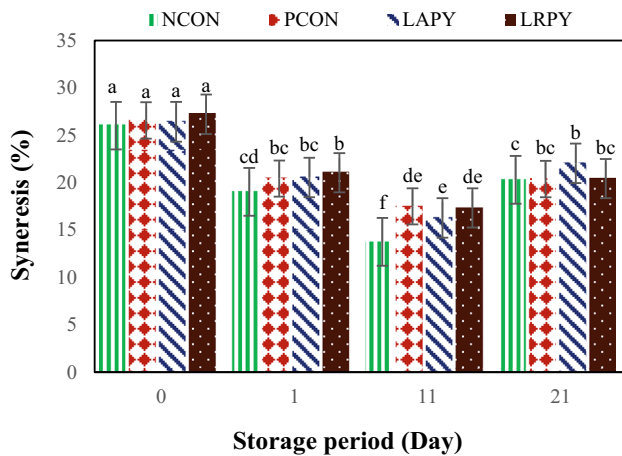


Fig. 3 Changes in the syneresis values of yogurt samples during cold storage. Different lowercase letters indicate significant differences at $p < 0.05$

(27.22%), while the lowest value was obtained in NCON sample (13.76%).

Furthermore, storage period had significant impact on the yogurt syneresis. The syneresis values of both controls and probiotic yogurts were meaningfully decreased ($p < 0.05$) until 11th day of the cold storage and thereafter, the amount of whey separation in yogurt samples increased noticeably. In agreement with our results, Tamjidi, et al. (2012) found that the separation of yogurt serum during storage had a decreasing trend from the first to 14th day of storage and thereafter it increased till the 21st day of storage. In contrary to our results, Yangilar and Çakmakçi (2017) and Güler-Akin and Akin (2007) reported a decrease in yogurt syneresis throughout cold storage period. As it is shown in Fig. 3, NCON samples had lower syneresis than PCON ones and this difference was significant at the 11th of storage period. The higher syneresis in PCON samples in comparison with NCON yogurts could be due to negative effect of added AFB₁ on metabolic activity of yogurt starter cultures, which cause an increase in net pressure in protein network (Akin, 1998).

Viability of LAB strains

Numerous parameters may affect the viability of lactic acid bacteria in yogurt including type of bacterial strain, inoculation level, pH, presence of hydrogen peroxide and dissolved oxygen, extent of produced metabolites such as lactic acid and acetic acids, concentration of solutes (osmotic pressure), buffering capacity of the media, storage and incubation temperature, storage and fermentation time, availability of nutrients as well as growth promoters and inhibitors (Donkor et al., 2006). The changes of viable cell count of lactic acid bacteria, from the preparation time of yogurt samples up to the end of cold storage are shown in

Fig. 4. There were significant differences ($p < 0.05$) among the viable counts of tested LAB strains; and in general, probiotics yogurt samples had higher bacterial counts than control samples. Although till the middle of storage period, LRPY yogurt samples had slightly the higher viable cell counts than LAPY samples, their number of viable cells at the end of storage were significantly reduced. As well, PCON samples had significantly lower bacterial counts than NCON and both probiotics yogurts throughout of storage period, indicated that AFB₁ had negative effect on starter cultures viability.

The adverse effect of aflatoxin M₁ on variability of LAB starter cultures have been previously demonstrated by Tajalli et al. (2014). Bacterial viability of all strains enhanced until 11th day of the storage and noticeably declined ($p < 0.01$) at the end of storage period. However, viable counts of both probiotic yogurts at the end of storage were above of standard limit ($> 10^7$ Log CFU/g). Similar findings were described by Birolo et al. (2000) and Güler-Akin and Akin (2007). However, in contrast with our results, Shaghghi et al. (2013) and Shah (2000a, b) observed no significant changes of bacterial cell count in commercial yogurts containing *L. acidophilus* and *B. bifidum* throughout the storage period.

AFB₁ binding ability of LAB strains

Generally, procedures to destroy the mycotoxins to the safe levels should have the following requirements: (1) deactivate or eliminate the toxin, (2) do not yield or release toxic remains in the foods and feeds, (3) sustain the nutritive values of the foods and feeds, (4) do not alter the sensory attributes and the quality characteristics of the product, and if feasible, (5) eliminate fungal spores (Park, 2002). Primary investigations have shown that the lactic acid bacteria

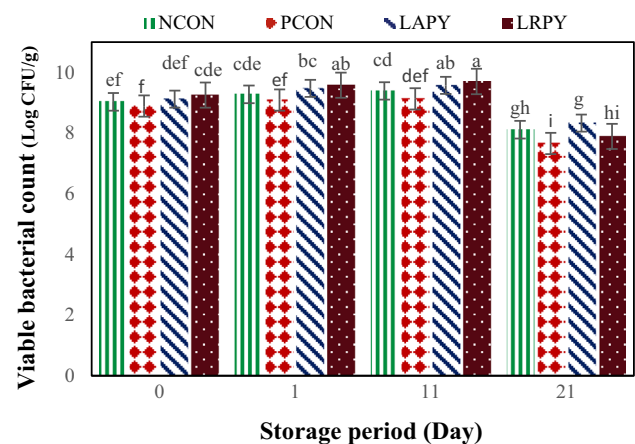


Fig. 4 Changes of viable cell counts of lactic acid bacteria during cold storage. Different lowercase letters indicate significant differences at $p < 0.05$

can remove aflatoxins, in vitro and in vivo models. Nevertheless, it is complicated to compare results of aflatoxin binding levels from various investigations, owing to the possible effect of technical variations.

Results in Fig. 5 show that all the strains tested were able to absorb AFB₁ efficiently, but at significant different levels ($p < 0.01$), which could be due to differences in the ability of their bacterial cell wall to absorb AFB₁. The amount of bound AFB₁ for all the yogurt treatments ranged from 64.56 ± 5.32 to $96.58 \pm 3.97\%$ during 21 days of the storage period. Elimination of AFB₁ by all tested lactic acid bacteria was rapid, since at least 64% of AFB₁ was removed after 1 day of storage. In agreement with our results, Tajalli et al. (2014) verified that more than 92% of AFM₁ was removed by *L. rhamnosus*. Elgerbi et al. (2006) also in an investigation on the capability of strains of *Lactobacillus* spp., *Lactococcus* spp. and *Bifidobacterium* spp. to bind AFM₁ declared that the extent of AFM₁ bounded after 96 h by these strains ranged from 4.5 to 73.1%. Besides, El-Nezami et al. (1998) represented 67% reduction of AFB₁ by *L. case* and Peltonen et al. (2001) reported 22.7–54.6% AFB₁ reduction for three strains of *L. rhamnosus*.

Among tested bacterial strains, probiotic yogurts were able to remove the higher level of AFB₁ in comparing with yogurt starter cultures, i.e. PCON sample throughout the storage period ($p < 0.01$). These findings were in accordance with the observations of Elsanhoty et al. (2014) who reported the higher AFM₁ reduction in probiotic yogurt contained an equal mixture of yogurt starter cultures and *L. plantrium* as compared to yogurt sample. As it is shown in Fig. 5, until the middle of storage period, LRPY probiotic yogurts had a little more binding capability to AFB₁, while at the end of storage, LAPY samples were able to eliminate slightly the higher percentages of AFB₁ ($p > 0.05$).

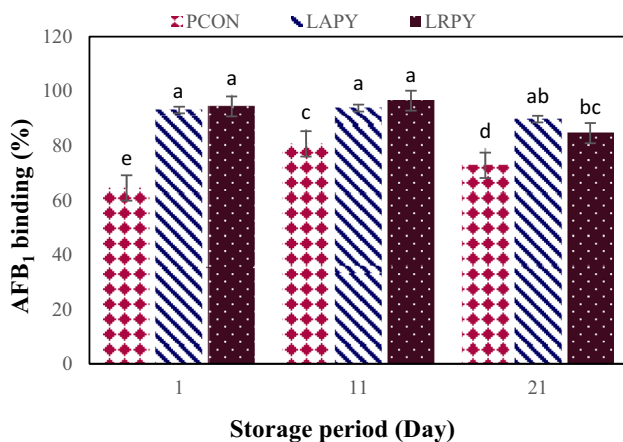


Fig. 5 Changes in the AFB₁ binding percentages by LAB in yogurt samples during cold storage. Different lowercase letters indicate significant differences at $p < 0.05$

Furthermore, the percentage of AFB₁ absorption was time dependent and significant interaction between treatment and storage period was found (Fig. 5). In general, by increasing the time of storage up to 11th day, the amount of AFB₁ binding ability by all yogurt samples significantly increased and reached to its maximum level but thereafter it noticeably decreased. The highest binding capacity of AFB₁ for PCON, LAPY and LRPY were recorded as $80.72 \pm 6.04\%$, $93.84 \pm 3.15\%$, $96.58 \pm 3.97\%$, respectively. As it demonstrated before (Fig. 4), the bacterial viability at the end of storage were significantly decreased by more than 1 log cycle and simultaneously substantial decrease of AFB₁ absorption ($p < 0.01$) were found at this period (Fig. 5). However, the reduction of AFB₁ binding in all the treatments throughout this storage interval were less than 10 percent. Similar results are reported by Abdelmotilib et al. (2018) and Tajalli et al. (2014). It is well confirmed that removal of AFB₁ depends neither on bacterial viability and nor on metabolic alteration of the toxin by bacteria. Indeed, toxin removal occurs by absorption to bacterial cell wall constituents rather than covalent linkage or metabolic deterioration, and dead cells still show binding capability (Haskard et al., 2000, 2001). It is even indicated that the bacterial cell wall disruption due to the heat processing result in the bacteria surface becomes more reachable to form extra AFB₁-bacteria linkage (Assaf et al., 2018; Liew et al., 2018). By bacterial absorption, toxin bioavailability is reduced, and consequently AFB₁ uptake and its entrance to systemic circulation are also constricted (Solis-Cruz et al., 2018).

Stability of the bacteria-AFB₁ complex during storage period

In addition to the kind of washing solution, the most important factors which affect the amount of complex stability (CS) of mycotoxins are the kind of bacteria strains and the level of toxin bound to bacterial cell. Bacteria strains differ due to the variations in their toxin-cell binding sites/cross-linked matrix that inhibits aflatoxin releasing (Utami et al., 2017). As well, the more aflatoxin adsorbed by bacterial cell, the longer time the adsorbed aflatoxin molecules would remain on the cell surface of bacteria (Lee et al., 2003).

The efficiency of selected LAB strains to bind AFB₁ after continual washing with PBS (pH ~ 7) is presented in Fig. 6. In the CS assay, significant differences were found between yogurt treatments in regarding to release of AFB₁ during 21 days of storage. The values of residual bound aflatoxin after three washes by bacteria during the storage time ranged from 73.25 ± 3.41 – $99.16 \pm 3.13\%$. Related to complexes stability, all of the treatments were shown different behaviors through the storage time. Initially, the

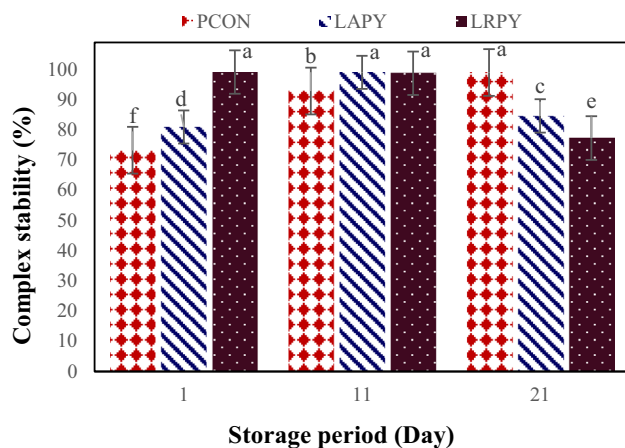


Fig. 6 Changes in the stability of the bacteria-AFB₁ complex in yogurt samples during cold storage. Different lowercase letters indicate significant differences at $p < 0.05$

CS of PCON was low (73.25%), but gradually increased to $92.88 \pm 3.59\%$ on the eleventh day and at the end of storage reached to its maximum level ($99.01 \pm 3.16\%$). The stability of the *L. acidophilus*-AFB₁ complex, at the beginning was $80.97 \pm 3.20\%$ and at 11th day of storage reached to its maximum level, i.e. $99.07 \pm 3.07\%$, and then lessened to $84.06 \pm 3.24\%$ at the end of product shelf life. Although LRPY contained *L. rhamnosus* showed the highest CS among the other treatments during the 21-days storage, it exhibited the lowest CS at the end of cold storage. Its highest CS at the beginning (99.16%), slightly decreased till 11th of storage day ($98.78 \pm 3.11\%$) and significantly reduced ($77.31 \pm 3.52\%$) after 21 days' storage, indicating a minimum stability among the examined samples. Besides, considering the amount of bound AFB₁ to the bacteria cells after washing (AFB₁-AW), there were significant variations ($p < 0.05$) amongst yogurt treatments and the values for PCON, LAPY and LRPY samples at the end of storage were $72.16 \pm 3.46\%$, $75.99 \pm 3.57\%$ and $65.44 \pm 3.61\%$, respectively. However, results showed that LRPY samples had significantly the higher AFB₁-AW than PCON (at 1st and 11th day of storage) and LAPY (at 1st day of storage) yogurts. Thus, in contrast to probiotic strains, yogurt starter cultures retained their AFB₁-AW at the end of storage period. The obvious superiority of the yogurt starter cultures in toxin elimination at the end of storage may be due to its acid toleration which retains their cell integrity and prevents release of toxin from the bacterial cell wall.

The data for CS-AFB₁ obtained at the initial and middle of storage are similar with those reported by Kabak and Var (2008) who reported 92.46–95.38% CS for AFM₁ by *Lactobacillus* and *Bifidobacterium* strains and Topcu et al. (2010) who reported a considerable CS for AFB₁ (77–83%) and patulin (75–81%) by *Enterococcus faecium* strains

after three washes with PBS solution. Hernandez-Mendoza et al. (2009) also found that after PBS solution, approximately 60–70% of AFB₁ binding to the bacterial cells was remained, proposing that the toxin is involved to the surface of bacteria. However, Elgerbi et al. (2006) contrarily stated that after the initial rinsing of LAB strains with PBS, the amount of AFM₁ liberated by bacterial cells was 85.7% (i.e. 14.3% CS) and after the third washing, nearly all adsorbed AFM₁ was released by the selected bacteria. Similarly, Shah and Wu (1999) revealed that only 10–40% of the bound AFB₁ was remained by probiotic strains when washed with water. Haskard et al. (2001) also reported a reversible binding of AFB₁ after five washes.

It should be considered that the variations in results may be elucidated by the alterations in extraction techniques, inconsistency in milk composition, method of milk contamination, concentration of mycotoxin, time elapsed before analysis, storage temperature, and characteristics of LAB strains utilized for yogurt production (Motawee and Abd El-Ghany, 2011).

An innovative biological method to decrease the health risks of mycotoxins via binding the toxins is application of probiotic strains of lactic acid bacteria. This work demonstrated that lactic acid producing bacteria have a great capacity to adsorb AFB₁ in yogurt. Our findings also revealed that binding of AFB₁ was an irreversible progress because after three washes with PBS, all tested bacteria liberated a small amount of bound AFB₁. Among tested LAB strains, probiotic yogurts particularly LRPY samples (yogurts containing *L. rhamnosus*) except at the end of storage had more AFB₁ binding capacity and more complex stability. Based on statistical results related to AFB₁-binding assay after washing the bacterial cells, no significant differences between PCON and LAPY samples at the end of storage were found, while LRPY samples (yogurts containing *L. rhamnosus*) had significantly lower AFB₁-binding assay after bacterial cells washing.

Mycotoxin-adsorbing bacteria should be capable to bind the toxins sturdily and must retain their cell integrity without dissociating. It is well documented that the most important factors for the selection of LAB as probiotic strain are their ability to tolerate the harsh gastrointestinal tract conditions, i.e. the toxicity of bile salts and gastric acid conditions. On the other hand, the ability of probiotic strain to resist against mucins and even their utilization as substrate is one of the key factors to be considered in the probiotic selection criteria (Kirjavainen et al., 1998; Salminen et al., 1996). These abilities enable probiotic microorganisms to persist in digestive tract and resulted in proper bacteria colonization and mycotoxin reduction. Therefore, consumption of probiotic yogurt is safer than non-probiotic ones; particularly in the regions where milk is considerably contaminated to mycotoxins. However,

further studies concerning the stability of the bacterial cells-AFB₁ complex specially under gastrointestinal tract condition are required and more investigations are needed to understand the various potential mechanisms underlying probiotic action.

Acknowledgements The authors acknowledge the financial support provided by Agricultural Sciences and Natural Resources University of Khuzestan.

Compliance with ethical standards

Conflict of interest The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

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