

Biodecontamination of milk and dairy products by probiotics: Boon for bane

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REVIEW ARTICLE

Abstract

In recent decades, “contamination of the environment, food, and feed by different contaminants such as heavy metals and toxins is increasing due to industrial life.” Commercial milk and milk products can be contaminated with heavy metals and mycotoxins. Biosorption is a low-cost method and has good potential for decontamination. In dairy products, “various starters, especially probiotics, can be used as biosorbants, while microorganisms are able to bind to heavy metals and toxins and decrease their bioavailability and hazards in the human body.” In this article, the key role of dairy starters and probiotics in the decontamination of toxins and heavy metals, and the best probiotics for decontamination of aflatoxins and heavy metals has been reviewed. After a quick glance at introducing dairy products and the main risks in association with the intake of some hazardous materials from dairy products, the application of biological systems is mentioned. Then, the article is focused on the role of beneficial microorganisms as the last chance to decrease the risk of exposure to toxins and heavy metals in dairy products. This review can be helpful for biotechnologists and scientists who have challenges about the existence of heavy metals and toxins in milk and dairy products, and help them to find the best method to decrease the content of the usual contaminants.

Keywords: aflatoxins; biosorption; decontamination; heavy metals; dairy products

Introduction

The World Health Organization (WHO) defines food safety as, “Approaches and methods for certifying the manufacture, maintenance, distribution and utilization of food happen in an assured system.” However, some people defined safe food as food without any contamination (El Sheikha, 2015).

Heavy metals naturally exist in the environment. Industrial activities can increase their content in air and soil, leading to phytotoxicity of plants (Asati *et al.*, 2016; Yang *et al.*, 2018). Milk and dairy products have an important role in the human food chain, especially children’s food; so, contamination of dairy products by toxins and heavy metals is one of the most important issues that can negatively impact consumers’ health. Milk and dairy

products can be contaminated with heavy metals under certain conditions through contamination of water and animal feed with environmental contaminants such as metal and cement smelters, sewage effluents, and industrial waste. Heavy metals' accumulation in milk can easily enter the human body and be dangerous for consumer's health (Abedi *et al.*, 2020). Dairy product contamination (heavy metal and aflatoxin) is very common all over the world (Ziarati *et al.*, 2018).

Heavy metals' toxicity occurs in levels of about 1.0–10 mg/L; however, lead and cadmium could have a toxic effect in 1–100 µg/L (Alkorta *et al.*, 2004). For example, different levels of exposure to cadmium could cause renal dysfunction, hepatic injury, and lung damage (Miura *et al.*, 2017; Naidoo *et al.*, 2019; Zhang *et al.*, 2014). Arsenic poisoning can cause death through disorder in essential metabolic enzymes (Khairul *et al.*, 2017). Maximum permissible limits of heavy metal contents in milk (considered by International Dairy Federation) are 2.6 µg/kg for cadmium, 10 µg/kg for Copper, 20 µg/kg for lead, and 328 µg/kg for zinc (Malhat *et al.*, 2012).

Aflatoxins directly (through eating contaminated food) and indirectly (primary contaminated products such as milk of contaminated livestock) can enter into the human body by the use of contaminated dairy products. Aflatoxins can cause negative effects on human health, such as liver or kidney cancer and chronic intoxications (Karazhiyan *et al.*, 2016). The most common aflatoxin in

dairy products is aflatoxin M₁ (AFM₁). AFM₁ is a metabolite of aflatoxin B₁ (AFB₁) after ingestion of contaminated feed (AFB₁) by livestock. About 0.3 to 6.2% of AFB₁ (Abdelmotilib *et al.*, 2018) can be bio-transformed into AFM₁ (4-hydroxy- AFB₁) and can excrete into milk and urine (Iha *et al.*, 2013; Karazhiyan *et al.*, 2016). AFM₁ is carcinogenic and toxicogenic, and can resist pasteurization and sterilization processes (Gonçalves *et al.*, 2020). AFM₁ compared with AFB₁ is approximately 10 times less mutagenic, genotoxic, and toxigenic. Its carcinogenic effects are displayed in different kinds of species (Elsanhoty *et al.*, 2014). AFM₁ can also cause gene mutation, DNA damage, cell transformation in mammalian cells, and chromosomal anomalies. Food and Drug Administration (FDA) and the European Commission recommended that the maximum permissible limits of AFM₁ in milk are 0.5 µg/kg and 0.05 µg/kg, respectively (Commission, 2006; FDA, 2019)

It is reported that mycotoxins in milk and dairy products, which can be produced by different kinds of fungi are: Aflatoxins (by *Aspergillus*), Compactin (by *Penicillium*), Cyclopaldic acid (by *Penicillium*), and Patulin (by *Penicillium*) (El Sheikha, 2019).

Many reports have investigated regarding contamination of milk by heavy metals and toxins all over the world. According to Tables 1 and 2, which present some of the above reports, the amount of lead in Iraq, Brazil, China, Spain, and Italy was more than the maximum permissible

Table 1. Some important data about milk contamination to heavy metals (from 2014 to 2021).

Country	Contamination	Concentration	Reference
Egypt	Pb	0.044–0.751 mg/L	Meshref <i>et al.</i> , 2014
	Cd	0.008–0.179 mg/L	
Serbia	Pb	54.3–95.2 lg/kg	Suturović <i>et al.</i> , 2014
	Cd	2.13–4.82 lg/kg	
Iraq	Pb	32 µg/L	Alani and Al-Azzawi, 2015
Pakistan	Pb	0.014 mg/Kg	Ismail <i>et al.</i> , 2015
	Cd	0.001 mg/Kg	
Bangladesh	Pb	0.2 mg/L	Muhib <i>et al.</i> , 2016
	Cd	0.073 mg/L	
Iran	Pb	14.0 µg/kg	Shahbazi <i>et al.</i> , 2016
	Cd	1.11 µg/kg	
Brazil	Pb	2.12–37.36µg/L	de Oliveira <i>et al.</i> , 2017
Mexico	Pb	0.03 mg/Kg	Castro-González <i>et al.</i> , 2018
	As	0.12 mg/Kg	
Poland	Pb	5.24 µg/L	Halagarda <i>et al.</i> , 2018
	Cd	0.15 µg/L	
Turkey	Pb	0.0055 mg/L	Seğmenoğlu and Baydan, 2021
	Cd	0.088 mg/L	
	As	0.002 mg/L	

As: Arsenic, Cd: Cadmium, Pb: Lead

Table 2. Some important data about milk contamination to mycotoxins in world from 2014 to 2021.

Country	Contamination	Concentration	Reference
Croatia	AFM ₁	0.003–1.135 µg/L	Bilandžić <i>et al.</i> , 2014
China	AFM ₁	80.4 ng/kg	Huang <i>et al.</i> , 2014
	OA	56.7 ng/kg	
	ZEA	14.9 ng/kg	
	α-ZEA	24.3 ng/kg	
Serbia	AFM ₁	0.01–1.2 µg/kg	Kos <i>et al.</i> , 2014
Iran	AFM ₁	> 0.05 µg/L	Fallah <i>et al.</i> , 2015
Macedonia	AFM ₁	408.1 ng/L	Dimitrieska-Stojković <i>et al.</i> , 2016
Pakistan	AFM ₁	>2610 ng/L	Aslam <i>et al.</i> , 2016
Argentina	AFM ₁	293 ng/L	Michlig <i>et al.</i> , 2016
Bosnia and Herzegovina	AFM ₁	60 ng/L	Bilandžić <i>et al.</i> , 2016
Italy	AFM ₁	52 ng/L	De Roma <i>et al.</i> , 2017
Tanzania	AFM ₁	0.627 ng/mL	Karczmarczyk <i>et al.</i> , 2017
Malaysia	AFM ₁	144 ng/L	Shuib <i>et al.</i> , 2017
Kosovo	AFM ₁	83 ng/L	Camaj <i>et al.</i> , 2018
El Salvador	AFM ₁	Approximately 100 ng/L	Peña-Rodas <i>et al.</i> , 2018
Turkey	AFM ₁	78.69 ng/L	Eker <i>et al.</i> , 2019
Ethiopia	AFM ₁	207 ng/L	Zakaria <i>et al.</i> , 2019
Kenya	AFM ₁	4563 ng/L	Kuboka <i>et al.</i> , 2019
Brazil	AFM ₁	45.18 ng/L	Venâncio <i>et al.</i> , 2019
Ecuador	AFM ₁	0.0774 µg/kg	Puga-Torres <i>et al.</i> , 2020
Spain	AFM ₁	0.009–1.36 µg/kg	Rodríguez-Blanco <i>et al.</i> , 2020
India	AFM ₁	1116 ng/L	Sharma <i>et al.</i> , 2020
Morocco	AFM ₁	4.46 ± 14.09 ng/L	Mannani <i>et al.</i> , 2021
Malawi	AFM ₁	0.551 µg/L	Njombwa <i>et al.</i> , 2021
	AFB ₁	0.61 µg/kg	
Spain	AFM ₁	12.6 ng/kg	Bervis <i>et al.</i> , 2021
	AFB ₁	0.61 µg/kg	

AFM₁: Aflatoxin M₁, AFB₁: Aflatoxin B₁, OA: Ochratoxin A, ZEA: Zearalenone, α-ZEA: α-zearalenone.

limits. Also, AFM₁ in China and India, and cadmium in Poland and Spain, were higher than permissible limits. This information confirms the importance of decontamination in milk and dairy products.

There are different methods for the decontamination of dairy products, such as physical, chemical (reverse osmosis, ion exchange, freeze concentration, and evaporation) (Patterson and Minear, 2013), and biological methods (using different biomaterials such as bacteria and yeasts biomass, plants, and seaweeds) (Abdelmotilib *et al.*, 2018; Hashim and Chu, 2004; Hayat *et al.*, 2017; Satyapal *et al.*, 2016; Sulaymon *et al.*, 2013; Vishnoi *et al.*, 2014). Adsorption is one of the most important decontamination strategies in dairy products (Giovati *et al.*, 2015; Massoud *et al.*, 2019; Milanowski *et al.*, 2017; Porova *et al.*, 2014). There are different biosorbents, such as “algae, plants, yeasts, fungi, and bacteria,” for the bioremoval of toxins and metals in fermented dairy products (e.g., kefir, kumis, yogurt, and doogh). Probiotic bacteria can also be used for this purpose. Fermented dairy

products are very popular, and they have a perfect taste (El Sheikha *et al.*, 2018; Yerlikaya, 2014). Probiotics can reduce contamination (heavy metals and aflatoxins) in fermented dairy products (Zoghi *et al.*, 2014). They are widely used for bioremoval of toxins (Massoud *et al.*, 2018; Zoghi *et al.*, 2017, 2019) as well as heavy metals (arsenic, mercury, lead, and cadmium) (Hadiani *et al.*, 2018, 2019; Khosravi-Darani *et al.*, 2019), heterocyclic aromatic amines (Khosravi-Darani *et al.*, 2019; Sarlak, 2020), and even pesticides (Wochner *et al.*, 2018).

In this article, reports about the influence of adding starters and probiotics into the formulation of dairy products on the bioremoval of contaminations such as toxins and heavy metals are reviewed.

Starters and Probiotics in Dairy Products

Food fermentation by microorganisms is one of the most economic and widely practiced methods for improving

texture, flavor, and functionality, and also for enhancing the shelf life of food products (Ray *et al.*, 2014; Salque *et al.*, 2013). The fermentation process can be carried out with starter cultures to certify consistency in commercial products by using familiar microorganisms with favorable traits, such as a high amount of acidification

via the manufacture of lactic acid and/or the sprinkling of secondary metabolites in the product matrix (Ryan *et al.*, 2015). Different starters have been used for producing various dairy products all around the world. Some of these products and their starters are mentioned in Table 3.

Table 3. Some fermented dairy products and related starters.

Fermented dairy products	Country/Region of origin	Starters	Reference
Acidophilus milk	—	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus casei</i> , <i>Bifidobacterium bifidum</i> , <i>Lactobacillus bulgaricus</i> , <i>Streptococcus thermophilus</i>	Raftaniamiri <i>et al.</i> , 2010
Buttermilk	Egypt and Ethiopia	(cultured buttermilk) Lactic acid bacteria (e.g., <i>Lactococcus</i> , <i>Lactobacillus</i> , <i>Streptococcus</i> , and <i>Leuconostoc</i>)	El Sheikh and Montet, 2014; Kumar <i>et al.</i> , 2015
Cheese	—	(cheddar cheese) Lactic acid bacteria starter culture (<i>Lactococcus lactis</i> ssp. <i>lactis</i> , <i>Lactococcus lactis</i> ssp. <i>cremoris</i> , and <i>Streptococcus salivarius</i> spp. <i>thermophilus</i>)	Ferreira and Viljoen, 2003
Matzoon	Armenia	Lactic acid bacteria	Macori and Cotter, 2018
Leben	Arab World	(Leben from camel milk) <i>Lactococcus lactis</i> , <i>Lactobacillus pentosus</i> , <i>Lactobacillus plantarum</i> , <i>Lactobacillus brevis</i> , and <i>Pediococcus pentosaceus</i>	Fguiiri <i>et al.</i> , 2013
Kishk	Arab World	Freeze-dried yogurt starter culture	Tamime <i>et al.</i> , 2000
Kumis	Central Asia Turkic countries Central Asia	<i>Lactococcus lactis</i> subsp. <i>lactis</i> , <i>Streptococcus thermophilus</i> , <i>Lactobacillus delbrueckii</i> subsp. <i>bulgaricus</i> , <i>Lactobacillus helveticus</i> , <i>Lactobacillus casei</i> subsp. <i>Pseudoplantarum</i> , and <i>Lactobacillus brevis</i> <i>Kluyveromyces marxianus</i> var. <i>lactis</i> , <i>Saccharomyces cerevisiae</i> , <i>Candida inconspicua</i> , and <i>Candida maris</i>	Simova <i>et al.</i> , 2002
Ymer	Denmark	<i>Streptococcus lactis</i> , <i>Streptococcus diacetilactis</i> ., <i>Streptococcus cremoris</i> , and <i>Leuconostoc citrovorum</i>	Poulsen, 1970
Kefir	Estonia, Hungary, Greece, Latvia, Romania, Slovakia, Bosnia and Herzegovina	<i>Lactobacilli</i> <i>Lactococcus</i> Acetic acid bacteria and yeast	Garrote <i>et al.</i> , 2001
Dahi	India	<i>Lactobacillus case</i> or <i>Lactobacillus acidophilus</i>	Yadav <i>et al.</i> , 2005
Mishti doi	India	<i>Streptococcus salivarius</i> ssp. <i>Thermophiles</i> , <i>Lactobacillus acidophilus</i> , <i>Lactobacillus delbrueckii</i> ssp. <i>bulgaricus</i> , <i>Lactobacillus acidophilus</i> , <i>Lactococcus lactis</i> ssp. <i>lactis</i> , <i>Saccharomyces cerevisiae</i>	Gupta <i>et al.</i> , 2000
Matsoni	Georgia	<i>Lactobacillus Streptococcus</i> , <i>Kluyveromyces marxianus</i> , <i>Candida famata</i> , <i>Saccharomyces cerevisiae</i> , <i>Lodderomyces elongisporus</i> , <i>Kluyveromyces lactis</i>	Bokulich <i>et al.</i> , 2015
Wara	Africa	<i>Lactobacillus</i> sp., <i>Leuconostoc</i> sp., <i>Pediococcus</i> sp., <i>Lactococcus</i> sp., yeasts	El Sheikh and Montet, 2014
Biruni	Sudan	Lactic acid bacteria	El Sheikh and Montet, 2014
Mish	Sudan and Egypt	Lactic acid bacteria	El Sheikh and Montet, 2014
Rob	Sudan	Lactic acid bacteria	El Sheikh and Montet, 2014
Doogh	Iran	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus rhamnosus</i> , <i>Lactobacillus casei</i> , <i>Bifidobacterium lactis</i>	Sarlak <i>et al.</i> , 2017
Yogurt	Serbia	<i>Streptococcus thermophilus</i> and <i>Lactobacillus bulgaricus</i>	Elsanhoty <i>et al.</i> , 2014
Clabber	United States	Starters like Kefir	Dyomina <i>et al.</i> , 2017

FAO (2001) defined probiotics as, “viable microorganisms, that while ingested in sufficient amounts, exert health benefits on the host (FAO/WHO, 2001).” The main beneficial effects of probiotics on human health include mucosal immunity support, decreasing lactose intolerance, preventing respiratory infections or diarrheas, feasible hypocholesterolemia effects, prevention of intestinal pathogens, inhibition of colon cancer or inflammatory bowel disease (Sanders *et al.*, 2014; Yu *et al.*, 2015).

The application of microorganisms, especially probiotics, recently has been investigated for their potential to heavy metals and aflatoxins reduction (Zoghi *et al.*, 2014). Most species known as probiotic bacteria are *Bifidobacterium* (*B.*), *Lactobacillus* (*L.*), *Bacillus*, and yeast *Saccharomyces* (*S.*) *cerevisiae*, and some strains of *Escherichia* (*E.*) *coli*. A practical taxonomy of nonpathogenic, fermentative, and nontoxic probiotic bacteria is lactic acid bacteria (LAB), which are used widely in food industries (Zoghi *et al.*, 2017). LAB usually have gram-positive cell walls, and peptidoglycan is their main cell wall structural component; teichoic acid, lipoteichoic acid, some neutral polysaccharides, and a proteinous S-layer are their minor components (Zoghi *et al.*, 2014).

Toxins' Bioremoval in Milk and Dairy Products

In recent decades, several scientific studies have been done regarding decontamination in dairy products, especially the biological decontamination method. Some of these researches are mentioned in Table 4.

El Khoury *et al.* (2011) investigated the application of LAB including *L. bulgaricus* and *Streptococcus thermophilus* on the reduction of AFM₁. They showed that using LAB is a potential method to decrease AFM₁ with the higher efficiency of *L. bulgaricus* compared to *Streptococcus thermophilus*. They also mentioned that the level of AFM₁, which is bound by LAB, enhanced with increasing the time of inoculation (El Khoury *et al.*, 2011). The binding ability of yogurt cultures was different. It is suggested that the difference in the binding ability of LAB is attributed to the difference in their cell-wall structure (Sarimehmetoğlu and Küplülü, 2004).

In addition to LAB, using *S. cerevisiae* is considered as an effective way for microbial detoxification (Karazhiyan *et al.*, 2016). A systematic review by Campagnollo *et al.* (2020) focused on parameters influencing the binding process of AFM₁ by yeast. The overall binding level of yeast was reported as 52.05%, in which the lowest binding capacity was related to the yeast extract peptone and the highest binding was associated with the ruminal fluid. Also, different factors, including temperature, yeast, pH, and the type of aflatoxin, have been mentioned as the

major parameters in the process of decontamination (Campagnollo *et al.*, 2020). Moreover, the effect of different treated *S. cerevisiae*, including heat, acid, and ultrasound treated, on the binding with AFM₁ was assessed by Karazhiyan *et al.* (2016). Among all treated yeasts, acid treatment had the most positive impact on yeast cells for improving their binding ability to aflatoxins which can be attributed to the release of monomers from polysaccharides under acidic conditions and their further changes into aldehydes after breaking down of glycosides linkages. After acid treatment, heat-treated yeasts showed the highest binding ability due to protein denaturation and Maillard reaction product formation, which caused an increase in the permeability of cell walls. Comparison between viable and unviable yeasts (heat, acid, and ultrasound treated) exhibited higher efficiency of unviable cells, which indicates that such treatments increase the binding capacity of yeasts (Karazhiyan *et al.*, 2016).

In a study performed by Taheur *et al.* (2017), a novel strategy for the reduction of mycotoxins using kefir grains was examined. The results showed that kefir microorganism grains could adsorb 82 to 100% of AFB₁, zearalenone, and ochratoxin A after cultivation in milk. The main strains that were able to adsorb mycotoxins were *L. kefir*, *Kazachstania servazzii*, and *Acetobacter syzygii*. The *L. kefir* KFLM3 was found to be the most active strain with an adsorption rate of 80 to 100% of the mycotoxins, and *K. servazzii* KFGY7 was found to retain higher mycotoxin than others after the desorption experiments. As a result, kefir consumption can assist in diminishing gastrointestinal absorption of mycotoxins and their toxic effects (Taheur *et al.*, 2017).

Heavy Metals' Bioremoval in Milk and Dairy Products

In Table 5, investigations regarding heavy metal bioremoval in milk and dairy products are illustrated.

In two different studies by Massoud *et al.* (2019, 2020a), application of *S. cerevisiae* to reduce the concentrations of lead and cadmium in milk was examined. The optimization process was also performed considering three factors including contact time, concentrations of biomass, and initial content of heavy metals (Massoud *et al.*, 2019, 2020a). Generally, the rate of removal of heavy metals increased with an increase in the biomass, contact time, and concentration of heavy metals. They concluded that optimized conditions for lead removal were obtained after 4 days (at the end of storage time) with the content of 22×10^8 CFU/mL of yeast and 70 µg/L of lead in milk (Massoud *et al.*, 2019). Similarly, the optimized process for cadmium bioremoval was achieved after 4 days with 80 µg/L of cadmium and 30×10^8 CFU/mL of *S. cerevisiae*

Table 4. Aflatoxin decontamination in milk and dairy products.

Product	Microorganism	Removal w/w%	Contaminant	Conditions	Reference
Milk	<i>Lactobacillus rhamnosus</i> (milk whey medium)	46.0%	AFB ₁	Optimal condition: 60 min in pH 3.0	Bovo <i>et al.</i> , 2014
Milk	Kefir starters 1. <i>L. acidophilus</i> , <i>Bifidobacterium</i> , & <i>Streptococcus thermophilus</i> (<i>thermophilic lactic</i> culture) 2. <i>Lactococcus lactis</i> subsp. <i>cremoris</i> , <i>Leuconostoc</i> , <i>Lactococcus lactis</i> subsp. <i>lactis biovar diacetylactis</i> , <i>Lactococcus lactis</i> , subsp. <i>lactis</i> , 3. <i>Debaryomyces hansenii</i> , <i>Kluyveromyces marxianus</i> subsp. <i>marxianus</i> ., yeast pool, Lactic acid bacteria pool	Full kefir starters 11.67–34.66% Yeast pool 65.33–68.89% LAB pool 65%	AFM ₁	Toxin Concentration: 150, 200, and 250 ng/L Temperature: 4 °C Time: 7 days	Kamyar and Movassaghazani, 2017
Milk	<i>Lactobacillus helveticus</i>	85%	AFM ₁	Time: 60 min	Ismail <i>et al.</i> , 2017
Milk	<i>Saccharomyces cerevisiae</i>	81.3%	AFM ₁	Time: 48 h	Foroughi <i>et al.</i> , 2018
Yogurt	A: <i>S. thermophilus</i> & <i>L. bulgaricus</i> B: 50% <i>S. thermophilus</i> & <i>L.</i> <i>bulgaricus</i> 50% <i>L. planetarum</i> C: 50% <i>S. thermophilus</i> and <i>L.</i> <i>bulgaricus</i> , 50% <i>L. acidophilus</i>	Treatment B: Highest reduction 31.5–87.8%	AFM ₁	Temperature: 5°C, Storage time: 1, 3, 5, and 7 days	Elsanhoty <i>et al.</i> , 2014
Yoghurt	<i>Lactobacillus</i> <i>acidophilus</i>	90%	AFM ₁	10 ⁸ CFU/ mL, Initial concentration of AFM ₁ :0.1, 0.5, 0.75 µg/L	Adibpour <i>et al.</i> , 2016
Yoghurt	<i>Saccharomyces cerevisiae</i>	76.46%	AFM ₁	Aflatoxin M ₁ : 100, 500, and 750 g/ M ₁ in 1, 7, 14, and 21 days, yeast treatments: heat, acid, and ultrasound	Karazhiyan <i>et al.</i> , 2016
Yoghurt	<i>Lactobacillus plantarum</i> , <i>Bifidobacterium animalis</i> , <i>Bifidobacterium bifidum</i>	Yogurt starters and <i>B. bifidum</i> , <i>B. animalis</i> (60.8%), Yogurt starters and <i>L. plantarum</i> , <i>B. Bifidum</i> 55.1%)	AFM ₁	Storage time: 1 or 10 days	Sevim <i>et al.</i> , 2019
Yogurt	<i>L. plantarum</i> , <i>B. animalis</i> , & <i>B.</i> <i>bifidum</i> , <i>L. plantarum</i>	49–60%	AFM ₁	Contact time: 4 h Temperature: 42°C	Sevim <i>et al.</i> , 2019
Kefir	<i>Lactobacillus casei</i> & kefir starter	88.17%	AFM ₁	Aflatoxin M, 500 pg, Kefir starters 2, 4, 6, 8, 10%, <i>L. casei</i> : 0.1, 0.3, 0.5, 0.7, 0.9 % in 48 h	Sani <i>et al.</i> , 2014
Kefir	Kefir-grains	96.8%	AFG ₁	Toxin concentration 5, 10, 15, 20, 25 ng/g, Kefir grain:5, 10, 20, 10, 25%, in 0, 2, 4, 6, 8 h, at 20, 30, 40, 50, 60°C	Ansari <i>et al.</i> , 2015

(continues)

Table 4. Continued

Product	Microorganism	Removal w/w%	Contaminant	Conditions	Reference
Kefir	Kefir grains: <i>Lactobacillus kefir</i> , <i>Kazachstania servazzii</i> , <i>Acetobacter syzygii</i>	82–100%	AFB ₁ , ZEA, OA	1 µg/MI mycotoxin, Kefir grains 10% w/v in 24 at 25°C	Taheur et al., 2017
UHT skim milk	Lactic acid bacteria (<i>Lactobacillus rhamnosus</i> , <i>Lactobacillus delbrueckii</i> spp. <i>Bulgarius</i> , <i>Bifidobacterium lactis</i>), <i>Saccharomyces cerevisiae</i>	LAB pool (30 min): 11.5 ± 2.3% LAB (60 min): 11.7 ± 4.4%, Saccharomyces: (30 min), 90.3 ± 0.3%, Saccharomyces: 60 min, 92.7 ± 0.7%	AFM ₁	0.5 ng AFM ₁ mL ⁻¹ , LAB pool: 10 ¹⁰ cells mL ⁻¹ Yeast: 10 ⁹ cells mL ⁻¹ Contact time: 30 min or 60 min	Corassin et al., 2013
Fermented milk drink	<i>Lactobacillus casei</i> Shirota	AFB ₁ -lys reduction: 82.37%	Serum AFB ₁ -lysine adduct	4-week intervention phases, (A): probiotic drinks 2 twice a day (B): placebo for 6, 8, or 10 weeks	Redzwan et al., 2016
Doogh	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus rhamnosus</i> , <i>Lactobacillus casei</i> , , <i>Bifidobacterium lactis</i>	Day 28, <i>Lactobacillus acidophilus</i> : 98.8 ± 1.3%	AFM ₁	0.500 ppb toxin, 1,14, or 28 days at 5 °C, <i>L. acidophilus</i> 9 log cfu/mL	Sarlak et al., 2017
Ergo fermented milk	<i>L. plantarum</i> , <i>Lactobacillus acidophilus</i> , <i>Lactobacillus brevis</i> , <i>Lactobacillus casei</i> subsp. <i>casei</i> , <i>Lactobacillus helveticus</i> , <i>Streptococcus faecalis</i> , <i>Streptococcus thermophiles</i> , <i>Leuconostoc mesenteroides</i> , subsp. <i>cremoris</i>	57.33% 54.04%	AFM ₁	Time: 1–5 days Temperature: 25°C	Shigute and Washe, 2018

AFM₁: Aflatoxin M₁, AFB₁: Aflatoxin B₁, OA: Ochratoxin A, ZEA: Zearalenone, AFG₁: Aflatoxin G₁.

Table 5. Heavy metals decontamination in milk and dairy products.

Product	Microorganism	Contaminant	Removal% W/W	Conditions	Reference
Milk	<i>Saccharomyces cerevisiae</i>	Pb	70%	Opt. at 22×10 ⁸ CFU inoculation of yeast, Lead content 70 µg/l	Massoud et al., 2019
Kefir	<i>Lactococcus lactis</i> , <i>Kluyveromyces marxianus</i> , co-culture	Ni, Cu, Cd, Pb, Fe	81.53%, 73.45%, 79.48%, 68.53%, 58.17%	Time: 10 days	Cherni et al., 2020
Milk	<i>Saccharomyces cerevisiae</i>	Cd	70%	Cadmium content in milk 80 µg/L, 30×10 ⁸ CFU <i>Saccharomyces cerevisiae</i> , storage time the 4th day,	Masoud et al., 2020
Milk	<i>Lactobacillus acidophilus</i>	Pb Cd	80% 75%	1 × 10 ¹² CFU of <i>L. acidophilus</i> , in 4 days with the initial pollution of 100 µg/L.	Massoud et al., 2020b
Milk	<i>Saccharomyces cerevisiae</i>	Hg	70%	Contact time: 30 days, initial concentration of Hg: 80 µg/L and biomass dosage 22 × 10 ⁸ CFU	Massoud et al., 2021

Lead: Pb, Nickel: Ni, Copper: Cu, Cadmium: Cd, Iron: Fe, Mercury: Hg.

(Massoud *et al.*, 2020a). Therefore, they have introduced applying *S. cerevisiae* as a novel and useful technology for the bioremoval of heavy metals from foodstuff (Massoud *et al.*, 2019, 2020a)

Different treatments, such as caustic, ethanol, acidic, and heat, can enhance the biosorption of heavy metals by microorganisms. In a study by Yekta Göksungur *et al.* (2005), “potential of baker’s yeast in bioremoval of cadmium and lead with 3 pretreatments (caustic, heat and ethanol)” was examined. Ethanol-treated yeast strains could remove the most content of metals and it can be explained by improving the availability of yeast binding sites and maybe enhancing the metals accessibility (Göksungur *et al.*, 2005).

Mechanisms of Bioremoval and Stability of Complexes (Probiotics/Starters-heavy Metal/Toxin)

AFM₁ and other toxins are accumulated in milk and dairy products because they are able to bind to milk protein components such as casein (Dyomina *et al.*, 2017; Granados-Chinchilla, 2016; Sarlak *et al.*, 2017). Therefore, numerous investigations have been focused on the removal of toxins using microorganisms, such as LAB (Dyomina *et al.*, 2017; Sarlak *et al.*, 2017).

Although the mechanism of bioremoval of toxins and heavy metals by LAB was not well known until now, it is proposed that toxins are highly linked by cell wall components of microorganisms and are not metabolically degraded (Zoghi *et al.*, 2014). Yeast and LAB are used widely to reduce toxins and metal ions. As both viable and dead cells are capable of adsorbing toxins, it is sensible to conclude that the removal of toxins is by adhesion to the components of microorganism’s cell wall relative to covalent binding, as reviewed by Shetty *et al.* (2006) (Shetty and Jespersen, 2006). It is indicated that mannan components of the *S. cerevisiae* cell wall play an important role in toxin binding (Devegowda *et al.*, 1996). Generally, the cell wall proteins of *S. cerevisiae* are bound to β -1,3-glucans by covalent linkage by β -1,6-glucan chains (Shetty and Jespersen, 2006). Apart from this, the major part of the LAB cell is made up of peptidoglycan, which contains teichoic and lipoteichoic acids. Also, a proteinous S-layer and neutral polysaccharides as components of the LAB cell wall have been recognized and reviewed by Lahtinen *et al.* (2004).

A study by Yiannikouris *et al.* (2004) indicated the interactions between zearalenone and β -D-glucans, in which β -1,3 D-glucan chains constitute a stable helical link with zearalenone and stabilized by β -1,6 D-glucan chains (Yiannikouris *et al.*, 2004). In order to investigate the

mechanism of binding of aflatoxins to *L. rhamnosus* it is indicated that carbohydrates in the cell wall are predominantly responsible for binding to aflatoxins. In samples treated by urea, it is shown that hydrophobic interactions play a significant role in binding, and treatment by NaCl and CaCl₂ showed that electrostatic interactions played a minor role (Haskard *et al.*, 2000).

Also, it is stated that AFM₁ is bound to LAB cell wall components by weak noncovalent interactions. The difference in the binding ability among different microorganisms is attributed to the cell wall and cell envelope structures (El Khoury *et al.*, 2011). Similarly, Turbic *et al.* (2002) mentioned that the different binding ability of LAB highly depended on the strain of the microorganisms (Turbic *et al.*, 2002).

Another study associated with the mechanism of biosorption illustrated that nonviable cells, including heat and acid-treated cells, produced complexes with higher stability, which means better access of groups in treated cells rather than viable ones. This phenomenon emphasizes that the viability of cells is not an important factor for the binding ability of cells (Haskard *et al.*, 2001). Furthermore, it is shown that acids might be capable of breaking amine binding in peptides and proteins, which leads to the production of peptides and even amino acids, and consequently, more accessible aflatoxin binding sites will be available (El-Nezami *et al.*, 2002). Similarly, it is noted that hydrophobic interactions are highly expected in LAB, which is treated by acid because acid treatment leads to denaturation of proteins and enhanced hydrophobic binding sites (Haskard *et al.*, 2000).

Moreover, the mechanisms of bioremoval could be influenced by various factors including types of microorganisms or even the status of biomass (living or nonliving microorganism), chemical properties of toxic materials, and environmental factors, such as temperature as well as pH (Javanbakht *et al.*, 2014).

For more illustration, Javanbakht *et al.* (2014) investigated the mechanism of removal of heavy metals by microorganisms. They suggested that two different types of pathways are involved in biosorption, which depends on cell metabolism and is divided into metabolism-dependent and metabolism-independent groups. The first pathway only occurs in viable cells through the transformation of metals across the cell wall. The second mechanism is involved in the physicochemical interaction between metals and functional groups of cell surface such as physical adsorption and ion exchange without depending on the cell metabolisms (Javanbakht *et al.*, 2014).

To investigate the stability of complexes, Haskard *et al.* (2001) evaluated the stability of 12 complexes between

LAB and AFB₁ considering both viable and nonviable cells and concluded that 71% of AFB₁ remained bound, indicating the high stability of the complexes. Also, they showed that nonviable cells retained a higher amount of AFB₁, as mentioned above (Haskard *et al.*, 2001). Based on their results, the stability of complexes depends upon three factors including strain, treatment type, and environmental conditions. Fazeli *et al.* (2009) conducted a study to investigate the effect of strains, including *L. casei*, *L. plantarum*, and *L. fermentum*, on the reduction of AFB₁ and concluded that all the strains were able to remove AFB₁, although *L. casei* was found to be a stronger binder of AFB₁ rather than other bacteria (Fazeli *et al.*, 2009).

A Study by Zoghi *et al.* (2020) showed that adsorption of patulin by LAB can be reversible in simulated gastrointestinal conditions. The reversibility of binding between LAB and patulin can be explained by the sense of non-covalent electrostatic bonds (Van der Waals and hydrogen bonds) (Zoghi *et al.*, 2020). Similarly, in another study, the adsorption of AFB₁, zearalenone, and ochratoxin A by kefir grains in simulated gastrointestinal pH was reversible. In pH 3, further amounts of toxins were released (Taheur *et al.*, 2017). Moreover, reduction of AFB₁ from a gastrointestinal model by several cells, including *L. rhamnosus*, *L. plantarum*, and *L. acidophilus*, were examined by Motameny *et al.* (2012), and they concluded that *L. plantarum* was the most active cell (Motameny *et al.*, 2012).

Conclusions

Aflatoxins and heavy metals frequently contaminate milk and dairy products at different levels. In the food industry, controlling aflatoxin and heavy metal levels in dairy products is a challenge for researchers. According to the recent studies summarized in this review, it is revealed that using different microorganisms (such as probiotics) in different dairy products could result in the removal of toxins and heavy metals by creating bonds between contaminants and these microorganisms. Using the starters in fermented dairy products can be helpful in the decontamination of toxins and heavy metals. According to this review, *L. bulgaricus*, Kefir grains, *L. acidophilus*, and *L. rhamnosus* could be useful for decreasing AFM₁ and other toxins in milk and dairy products. Also, for decontamination of heavy metals, kefir grains had the best ability for the bioremoval of different metals.

Future directions

More investigations are needed regarding the stability of binding between probiotics and toxins/heavy metals in

in vivo and *in vitro* conditions. Also, more experiments should be done for finding optimum conditions for special starters in special dairy products for better decontamination.

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RM was involved in writing and original draft preparation; AZ was responsible for writing, review, and editing; KKD was concerned with conceptualization and supervision; FM was involved in writing and editing; and SJ, RM, and YR were responsible for review and editing.

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