










Microbiological quality of home production of sprouts in Brazil

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Abstract

Sprouts are considered foods with high nutritional value, tasty, and easy to produce even in domestic environments. However, their consumption has been linked to various food outbreaks, as they are susceptible to microbial contamination. This work aimed to evaluate the microbiological quality of lentil, wheat, and chia sprouts germinated in a domestic environment and the effectiveness of the sprout and seed sanitization process. Total coliforms, *Escherichia coli*, mesophilic aerobes, and *Bacillus cereus* were determined, and the presence of *Salmonella* spp. and *Listeria monocytogenes* was investigated. No strains of *Salmonella* spp., *E. coli*, and *L. monocytogenes* were found. However, a high microbial load of total coliforms was found in the sprouts. *B. cereus* was found only in the wheat and chia sprout samples. The sanitization process with sodium hypochlorite was able to reduce the microbial load of the sprouts, but the load remained high. Seeds should be marketed only when they are suitable for home germination after going through a more efficient decontamination process. In addition, competent authorities are supposed to create and widely disseminate a document of good production practices for safer domestic cultivation of sprouts.

Keywords: bacteria; germination; seed disinfection; total coliforms; chemical decontamination.

Practical Application: Sprouts produced in a domestic environment showed a high microbial load, even after sanitization with sodium hypochlorite.

1 INTRODUCTION

Some countries, such as Brazil, the United States, and Australia, have nutritional guidelines that advocate that the consumption of fresh or minimally processed foods should be the basis for a nutritionally balanced, tasty, and sustainable diet, and this group includes foods such as grains, tubers, roots, vegetables, and fruits (Brasil, 2014; Australia, 2013; United States of America [USA], 2020). Owing to an increase in the number of individuals with chronic non-communicable diseases over the years, consumption of these foods is increasingly important, since a plant-based diet is a preventive factor for these diseases (Barros et al., 2021).

However, people's current lifestyle, characterized by a lack of time, is one of the main reasons for an inadequate diet (Menezes & Maldonado, 2015). Although cooking requires time and planning, it can contribute to better diet quality. Preparing and cooking one's own food helps the transition to and maintenance of a healthier lifestyle (Brasil, 2014; Oliveira & Castro, 2022).

Grains are staple foods that are widespread and consumed worldwide; they have an optimal nutritional composition and bring beneficial health effects (Aloo et al., 2021). Consumption

of sprouts has become popular worldwide because consumers have been searching for foods with high nutrient content. The sprouting process leads to several changes in the nutritional composition of grains, e.g., it improves digestibility, increases nutritional value, reduces antinutritional factors, and favors the accessibility of nutrients (Miyahira et al., 2021). Peñas & Martínez-Villaluenga (2020) stated that germination reactivates seed metabolism, inducing the degradation of macronutrients and antinutritional compounds, and is also able to form bioactive compounds that are beneficial to health. Salgado et al. (2022) reported that an increase in protein content, dietary fiber, and phenolic compounds and a decrease in antinutritional factors occurred after germination in chia sprouts, for example (Salgado et al., 2022). In wheat sprouts, germination can increase antioxidant activity (Miyahira et al., 2022). In lentils, this process can increase the concentration of protein and nutrients such as iron, zinc, and manganese (Santos et al., 2020). Thus, the cultivation of sprouts in the home environment does not require a large space, and it consists of simple, economical, and sustainable steps. Moreover, it is an excellent strategy to encourage consumers to participate in the entire process of production of this highly nutritious food until the moment it is actually available for consumption (Ebert, 2022).

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Despite the demonstrated benefits of grain germination, the conditions of this process are favorable to the proliferation of pathogenic microorganisms present in the grain. Increased moisture, ambient temperature, and high nutrient availability result in an ideal scenario for microbial growth (Brankatschk et al., 2014; Ding et al., 2013; Iacumin & Comi, 2019). Foodborne outbreaks have been associated with sprout consumption in different parts of the world between 1988 and 2020, totaling about 15,342 cases, 313 hospitalizations, and 61 deaths (Carstens et al., 2019; Miyahira & Antunes, 2021).

In order to minimize the risks involved in sprout production, the U.S. Food and Drug Administration (FDA), along with the Institute for Food Safety and Health (IFSH), created the Sprout Safety Alliance (SSA), a public–private alliance that develops training programs and implementation of best practices for safe sprout production (SSA/FDA). Proper decontamination treatments should inactivate pathogenic microorganisms while preserving seed viability, germination, and vigor (National Advisory Committee on Microbiological Criteria for Foods [NACMCF], 1999). Although complete elimination of bacterial contamination by seed disinfection treatments in sprout production is very difficult to achieve (European Food Safety Authority [EFSA], 2011), studies have evaluated different methods (chemical, biological, and physical methods) of seed disinfection and the impact of these processes on germination capacity (Ding et al., 2013; Sikin et al., 2013). According to Gilbert et al. (2023), heat treatments that significantly reduce microbial growth also decrease seed germination capacity and are therefore not eligible for seed disinfection. Among the various chemical sanitizers recommended, sodium hypochlorite and hypochlorous acid were the most effective in reducing bacterial load by at least five logarithmic units without adversely affecting seed germination (Gilbert et al., 2023).

Considering that germination increases the nutritional value of grains, and germination conditions may be conducive to the development of pathogenic microorganisms, the microbiological contamination of sprouts and the efficiency of chemical sanitization of seeds and sprouts need to be evaluated for further development of actions to support regulations that ensure the safe production of sprouts. The reason is that sprouts are being increasingly grown in domestic environments and used in salads or food preparations that do not undergo any heat treatment for pathogen reduction. Thus, this study aimed to evaluate the microbiological quality of lentil, wheat, and chia sprouts produced in the home environment, as well as the effectiveness of the sanitization process with sodium hypochlorite on the sprouts and seeds.

1.1 Relevance of the work

Sprouting is a simple method that can be carried out at home, so this study aimed to evaluate the microbiological quality of sprouts produced in a domestic environment and the effectiveness of the sanitization process with sodium hypochlorite. High levels of total coliforms and mesophilic aerobic microorganisms were found. The microbial load remained high after sanitization with sodium hypochlorite. The results suggest the need to develop strategies for the safe production of sprouts,

such as the marketing of seeds that have been treated and are suitable for germination and the dissemination of good sprout germination practices for home production.

2 MATERIALS AND METHODS

2.1 Samples and germination/sanitization process

Lentil (*Lens culinaris* L.), wheat (*Triticum aestivum* L.), and chia (*Salvia hispanica* L.) were divided into sanitized seed (SS) and unsanitized seed (USS) groups. After germination, the SS and USS groups were subdivided into four groups: unsanitized seed and unsanitized sprout (USS/USP), unsanitized seed and sanitized sprout (USS/SSP), sanitized seed and unsanitized sprout (SS/USP), and sanitized seed and sanitized sprout (SS/SSP). Figure 1 shows a sanitization/germination flowchart. The germination process was repeated five times for each of the four seed/sprout treatment groups, totaling 60 samples: 20 of lentil, 20 of wheat, and 20 of chia sprouts. Germination of lentil and wheat grains was performed according to Miyahira et al. (2022). The germination process of chia seeds was performed according to Abdel-Aty et al. (2021).

The sanitization process performed before (on the seeds) and after germination (on the sprouts) was the same. A solution of water (1 L) added to 10 mL of sodium hypochlorite at a concentration of 2% was prepared in a sterile container for 15 min (200 ppm), as specified on the product label by the manufacturer. Microbiological analyses were performed on the sprouts from March to October 2022, immediately after seed germination.

2.2 Microbiological analysis

2.2.1 Determination of total coliforms, *Escherichia coli*, mesophilic aerobic bacteria, and *Bacillus cereus*

Notably, 10 g of the sample was homogenized with 90 mL of 0.1% peptone water (10^{-1} dilution). Next, serial dilutions were made with 1 mL of the 10^{-1} dilution into a tube with 9 mL of 0.1% peptone water, performing the 10^{-2} dilution up to the 10^{-7} dilution.

Notably, 10 g of the sample was homogenized with 90 mL of 0.1% peptone water (10^{-1} dilution). Next, serial dilutions were made up to the 10^{-7} dilution (American Public Health Association [APHA], 2015). Total coliform and *E. coli* analyses were carried out using 3MTM Petrifilm™ plates. For mesophilic aerobic bacteria analysis, the pour plate technique was used, with plating of the dilutions on Standard Counting Agar (SCA) culture medium and incubation at $35^{\circ}\text{C} \pm 2^{\circ}\text{C}$ for 48 h (APHA, 2015). For the determination of *B. cereus*, the spread plate technique was used as described by International Organization for Standardization (ISO) method 7932 (ISO, 2004).

2.2.2 Detection of *Listeria monocytogenes*

Notably, 25 g of the sample was homogenized in 225 mL of Buffered Listeria Enrichment Broth (BLEB) and incubated at $30^{\circ}\text{C}/4$ h (Hitchins et al., 2022). Selective agents were added

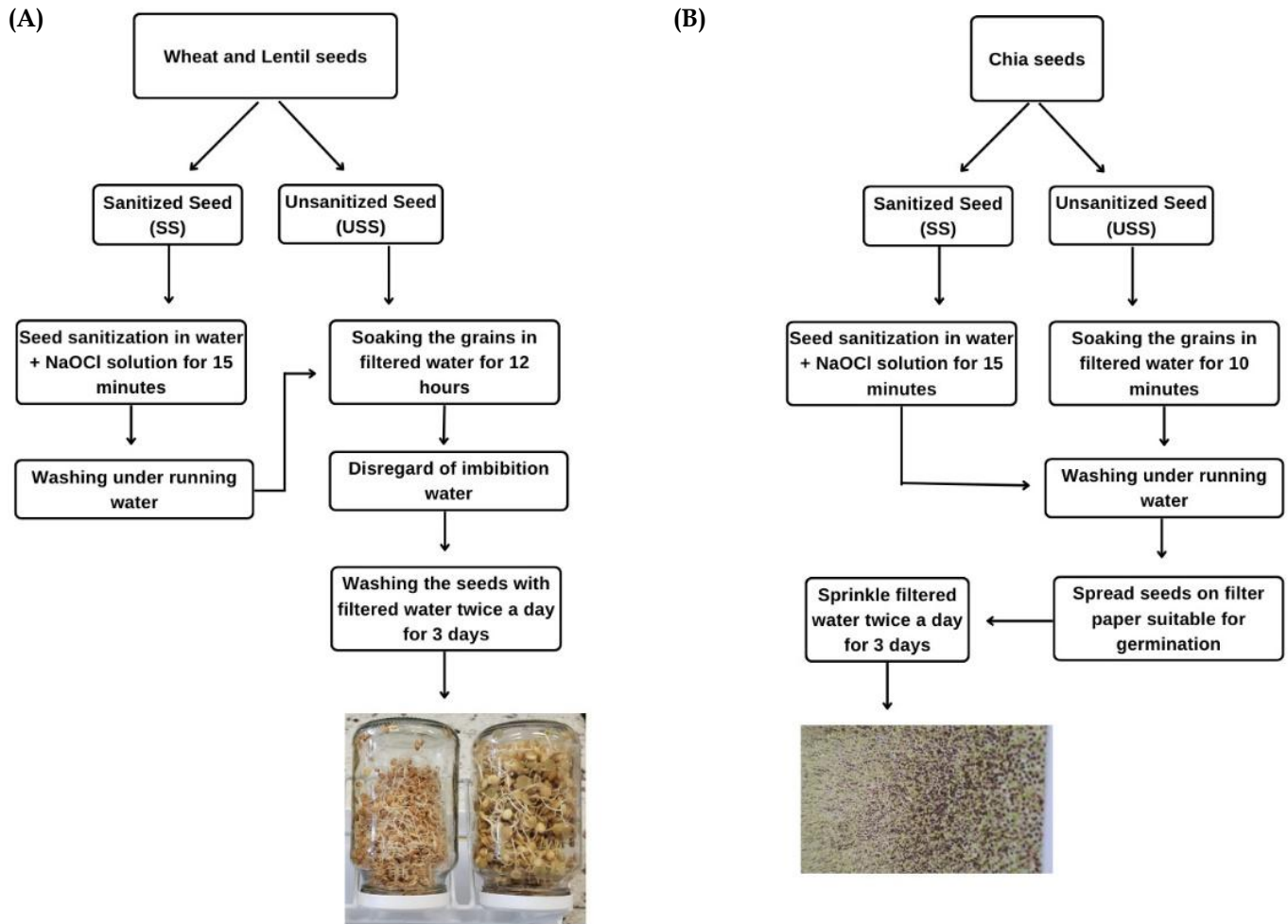


Figure 1. Flowchart of sanitization/germination of lenticil, wheat (A), and chia (B) seeds.

and incubated at 30°C/44 h. After incubation, one well from the selective enrichment vial was streaked onto an Oxford Agar plate, and the plates were incubated at 35°C/24–48 h. When present, typical colonies were streaked on Soybean Trypticase Agar plates supplemented with 0.6% yeast extract and incubated at 30°C/24–48 h. For confirmation, the catalase, motility, and hemolysis tests were performed

2.2.3 Detection of *Salmonella* spp.

Notably, 25 g of the sample was homogenized in 225 mL of 1% peptone water and incubated at 37°C/24 h. Then, 1 mL of the pre-enriched sample was transferred to tubes with the selective media Tetrathionate Broth and Selenite Cystine Broth and incubated at 37°C/24 h. After that, one batch of each broth was streaked onto Hecktoen's Enteric Agar and Xylose Lysine Agar plates and incubated at 37°C/24 h. Typical colonies were inoculated into tubes slanted with Lysine Iron Agar and Triple Iron Sugar Agar for confirmation (Andrews & Hamamack, 2007).

2.3 MALDI-TOF

Typical isolated *Salmonella* spp. and *L. monocytogenes* colonies were identified by mass spectrometry (matrix-assisted laser

desorption/ionization time-of-flight [MALDI-TOF]), according to the standard extraction protocol, using formic acid (Bruker, 2015), at the Medical Microbiology Research Laboratory of the Paulo de Góes Microbiology Institute at the Federal University of Rio de Janeiro.

2.4 Data analysis

The data collected from the microbiological analysis were analyzed and expressed in log CFU/g for total coliforms, *E. coli*, *B. cereus*, and total mesophilic aerobic bacteria, and in the presence/absence of *Salmonella* spp. and *L. monocytogenes* in 25 g of the sample. The results were presented as means and standard deviations calculated by Microsoft Excel. Analysis of variance (ANOVA) and Tukey's test were applied to determine statistically significant differences between the types of sanitization treatment, with a significance level of 5% ($p < .05$), using statistical software GraphPad Prism 9.

3 RESULTS AND DISCUSSION

The germination process is favorable to microbial growth, and some outbreaks involving sprouts have been reported in different parts of the world. The most recent were reported in

the year 2020, with 51 cases, with three hospitalizations, related to *E. coli* involving clover sprouts (Centers for Disease Control and Prevention [CDC], 2020), and in the year 2022, 63 people from eight states in the United States were infected with *Salmonella* Typhimurium strain after consuming raw alfalfa sprouts. It is estimated that the number of people infected in this latest outbreak was probably higher than reported because people recovered without medical care and without being tested for *Salmonella* (CDC, 2023).

A recent study showed that biohazard is not one of the main topics that consumers relate to sprouts. The study evaluated the perception of 315 Brazilian consumers about sprouts and showed that in a word association questionnaire, the “biological risk” was the least mentioned category. This result shows that few people related the consumption of sprouts with a negative aspect, such as foodborne diseases (FBD). This result shows that few people related the consumption of sprouts with a negative aspect, such as FBD (Miyahira et al., 2023). It is particularly relevant to assess the microbiological quality in the sprout production process to ensure consumption of a safe food, because dried seeds contain between 10^3 and 10^5 nonpathogenic bacteria per gram, which multiply rapidly during germination, and sprouts can contain between 10^8 and 10^9 nonpathogenic bacteria per gram until consumption (NACMCF, 1999). This very high bacterial population can make it difficult to detect pathogenic bacteria in sprouts. The location of bacteria on the surface of seeds/sprouts, their internalization potential, and their ability to adhere to seed/sprout tissues are also important factors that will influence the effectiveness of intervention strategies such as washing and decontamination (EFSA, 2011).

3.1 Total coliforms

In the present study, the results for total coliforms, in the lentil samples in which the sprouts were sanitized (USS/SSP), show that the values found were significantly lower than those in which the sprouts were not sanitized (USS/USP) (Table 1). In wheat, it was found that in the samples in which the sprouts had not been sanitized (USS/USP and SS/USP), there was a significantly higher microbial load of total coliforms than those that had been sanitized (USS/SSP and SS/SSP) (Table 1). For chia, it was found that the microbial load of total coliforms was significantly lower in the samples in which seeds and sprouts had been sanitized when compared to the samples in which the sprouts had not been sanitized, regardless of seed sanitization. It can be inferred that sanitization of the sprouts significantly reduced the number of total coliforms in the three types of study grains; however, it was found that the microbial load remained high, with values higher than 6.57 ± 1.65 log CFU/g. It is worth noting that when only the seed was sanitized, there was no significant reduction in the number of total coliforms in the three study sprouts.

Martínez-Villaluenga et al. (2008) conducted a study to evaluate the microbiological quality of broccoli and radish sprouts germinated for up to 5 days. Prior to germination, seeds were sanitized with 7% sodium hypochlorite for 30 min, washed and soaked with distilled water, and then placed in a seed germinator. Total coliforms were investigated, and an average value of 9.50 log CFU/g was found in broccoli seeds after germination. In the radish seeds, an average value of 8.04 log CFU/g was found. The values were similar to the one found in the present study, which is indicative of a high microbial load in the sprouts.

Table 1. Results of microbiological analyses of lentil, wheat, and chia sprouts before and after the seed and sprout sanitization process.

Sample	Microorganisms	USS/USP	USS/SSP	SS/USP	SS/SSP
		Mean \pm SD			
Lentil	Total coliform (log CFU/g)	8.55 \pm 0.09 ^a	7.99 \pm 0.32 ^b	8.63 \pm 0.02 ^a	7.32 \pm 0.13 ^c
	<i>Escherichia coli</i> (log CFU/g)	< 1	< 1	< 1	< 1
	Aerobic mesophilic (log CFU/g)	8.88 \pm 0.09 ^b	8.17 \pm 0.24 ^c	9.38 \pm 0.09 ^a	8.41 \pm 0.12 ^c
	<i>Bacillus cereus</i> (log CFU/g)	< 2	< 2	< 2	< 2
	<i>Listeria monocytogenes</i> /25 g	Absence	Absence	Absence	Absence
	<i>Salmonella</i> spp./25 g	Absence	Absence	Absence	Absence
Wheat	Total coliform (log CFU/g)	8.41 \pm 0.30 ^{ab}	7.84 \pm 0.05 ^c	8.60 \pm 0.31 ^a	8.07 \pm 0.12 ^{bc}
	<i>Escherichia coli</i> (log CFU/g)	< 1	< 1	< 1	< 1
	Aerobic mesophilic (log CFU/g)	9.12 \pm 0.10 ^a	8.74 \pm 0.25 ^a	8.64 \pm 0.41 ^a	7.93 \pm 0.48 ^b
	<i>Bacillus cereus</i> (log CFU/g)	5.46 \pm 0.51 ^{ab}	4.88 \pm 0.51 ^{ab}	5.61 \pm 0.62 ^a	4.65 \pm 0.47 ^b
	<i>Listeria monocytogenes</i> /25 g	Absence	Absence	Absence	Absence
	<i>Salmonella</i> spp./25 g	Absence	Absence	Absence	Absence
Chia	Total coliform (log CFU/g)	8.51 \pm 0.61 ^a	7.59 \pm 0.30 ^{ab}	8.46 \pm 0.14 ^{ab}	6.58 \pm 1.66 ^b
	<i>Escherichia coli</i> (log CFU/g)	< 1	< 1	< 1	< 1
	Aerobic mesophilic (log CFU/g)	9.21 \pm 0.26 ^a	7.87 \pm 0.71 ^{ab}	8.73 \pm 0.16 ^a	6.58 \pm 2.28 ^b
	<i>Bacillus cereus</i> (log CFU/g)	6.03 \pm 1.08 ^a	5.90 \pm 1.03 ^a	5.71 \pm 0.67 ^{ab}	4.22 \pm 0.29 ^b
	<i>Listeria monocytogenes</i> /25 g	Absence	Absence	Absence	Absence
	<i>Salmonella</i> spp./25 g	Absence	Absence	Absence	Absence

Caption: USS: unsanitized seed; SS: sanitized seed; USP: unsanitized sprout; SSP: sanitized sprout; SD: standard deviation; CFU: colony forming units; g: gram.

Identical letters in the same row indicate that there was no significant difference ($p < .05$) in the samples.

Another study by Tornuk et al. (2011) determined the number of total coliforms in wheat seeds and sprouts germinated for 9 days under different conditions of relative humidity (RH) (90 and 95%) and temperatures (18°C, 20°C, and 22°C) and investigated the disinfection capacity of sodium hypochlorite (NaOCl) (100, 200, and 400 ppm) and hydrogen peroxide (H₂O₂) (3 and 6%). The authors found a significant increase in total coliforms after germination and reported that increasing concentrations of NaOCl and H₂O₂ resulted in reductions in the total coliform population; there were greater reductions when seeds had been soaked in 400 ppm NaOCl for 30 min and then germinated at 18°C and 90% RH. It is worth noting that the authors also found that sodium hypochlorite at 200 ppm was also able to reduce total coliforms significantly by 0.54 log CFU/g. This result was different from that of the present study, which found no significant difference in the number of total coliforms after sanitizing only the seeds.

3.2 *Escherichia coli*

In this study, the presence of characteristic *E. coli* colonies was not found in any of the study samples. For this reason, it was not possible to evaluate the effectiveness of the sanitization process in reducing the amount of *E. coli* in the samples of this study. However, Tornuk et al. (2011), when analyzing the microbial load of wheat seeds and its changes during germination, found that there was an increase in the number of *E. coli* during germination in different conditions of humidity and temperature, reaching up to 6.37 log CFU/g at 90% humidity and 22°C temperature and 7.17 log CFU/g at 95% humidity and 22°C temperature. The authors also found that sanitization with sodium hypochlorite or hydrogen peroxide at different concentrations was able to significantly reduce the presence of this microorganism.

Jeddi et al. (2014) performed microbial evaluation of vegetables and sprouts of wheat and mung beans bagged in a supermarket chain and found the presence of *E. coli* in 12 of 64 sprout samples, corresponding to 18.7% of the study samples. Kim and Cheig (2021) evaluated the microbiological contamination of minimally processed products in Korea, and they found an average of 1 log CFU/g of *E. coli* in seven samples of mixed sprouts, which was considered to be within the acceptable limit according to the Korea Food Code standards. A literature review by Sikin et al. (2013) showed that several treatments are effective in decreasing the microbiological load of *E. coli* in seeds and sprouts; however, some sanitizers, such as organic acids, can impair the germination process or its sensory characteristics.

3.3 *Aerobic mesophilic bacteria*

The results of aerobic mesophilic (AM) in the lentil samples showed that the sanitized sprouts had statistically lower values than the unsanitized ones (Table 1), regardless of seed sanitization. Therefore, there was no reduction in the amount of AM when the seed was sanitized before sprouting, and the potential for AM internalization in lentil seeds and their structure may have influenced this result (EFSA, 2011). By contrast, in wheat, only SS/SSP had a significant reduction of AM when compared

to the other samples (Table 1); i.e., the use of sanitization, only when performed simultaneously on the seeds and the sprouts, was effective in reducing the microbial load in wheat sprouts. In chia, sprout sanitization significantly reduced the amount of AM, while sanitization of the seeds alone was not able to reduce AM significantly (Table 1).

Saroj et al. (2006) reported that vegetable seeds can contain less than 2 log CFU/g of mesophilic aerobic bacteria and that this population of microorganisms can increase rapidly during germination owing to favorable conditions for bacterial growth. The mean AM count found in the present study was similar to the one found by Seow et al. (2012), who reported mean AM counts in bean sprouts sold in Singapore of 8.0 log CFU/g. The same occurred with the study conducted by Tango et al. (2018); the authors found the value of 9.35 log CFU/g of AM in mixed sprouts sold in Korea that had not been disinfected for microorganism reduction. Tornuk et al. (2011) compared the effectiveness of chemical sanitization in reducing AM and found that treating wheat seeds with 400 ppm NaOCl reduced AM count by 0.29 log compared to the control, and further found a 1.46 log reduction in sprouts after seed sanitization when compared to sprouts without seed sanitization. The reduction in the amount of AM in the study of Tornuk et al. (2011) was much greater than that found in wheat in the present study, which was 0.48 log between USS/USP and SS/USP samples. The authors explained that chlorine-based sanitizers affect microbial cells in various ways, e.g., by increasing membrane permeability, inhibiting enzyme systems, and irreversibly damaging the DNA of bacterial cells.

3.4 *Bacillus cereus*

Bacillus cereus (*B. cereus*) was isolated only in the wheat and chia sprouts because this bacterium is more present in cereals (Rahnama et al., 2023). The results (Table 1) showed that the SS/SSP wheat sample had a significant reduction of *B. cereus* only when compared to SS/USP. Thus, it can be suggested that sanitization of both seeds and sprouts is not able to reduce the microbial load of *B. cereus* adequately. In comparison, SS/SSP chia samples showed a significant reduction of *B. cereus* only when compared to USS/USP samples, suggesting that sanitization of the seeds only or the sprouts alone is not able to reduce contamination by *B. cereus*.

It is worth noting that this microorganism is involved in outbreaks of foodborne illnesses associated with vegetable products (Choi & Kim, 2020; Glasset et al., 2016; Ultee et al., 1999) and sprouts (Portnoy et al., 1976). Foodborne illnesses caused by *B. cereus* usually occur if the product has more than 5 log CFU/g (Harmon et al., 1987). Although the number of *B. cereus* cells required to produce sufficient emetic toxin to cause diseases is still unclear, studies have shown that levels of 10³ to 10¹⁰ CFU/g have been found in foods involved in cases of emetic disease, and in most cases, there were at least 10⁵ CFU/g (Arnesen et al., 2007; Dietrich et al., 2021). Thus, the unsanitized wheat and chia sprouts in the present study could be a risk for the development of FBD, as they showed values greater than 5 log CFU/g. In addition, the values of *B. cereus* found in the wheat and chia sprouts in the present study are above the criteria established

by the Brazilian legislation (Brasil, 2022) of 5×10^2 CFU/g (2.69 log CFU/g); therefore, wheat sprouts can be rated as unfit for consumption in Brazil.

Just like this study, the one by Pao et al. (2005) examined the growth of *Bacillus* spp. in sprouts grown in a domestic environment and found no growth of *B. cereus* during germination in lentil, alfalfa, and mung bean sprouts. However, in radish and broccoli sprouts, they found values close to 5 log CFU/g. In a study conducted in natural products stores in the Washington area, *B. cereus* was found to have values ≥ 3 MPN/g in 69% of 98 seed samples tested (

Harmon et al., 1987), with the average microbial load being 5.39 log CFU/g for wheat sprouts, as found in the present study.

3.5 *Listeria monocytogenes*

Although *L. monocytogenes* has been related to some food outbreaks with sprouts (CDC, 2015; Garner & Kathariou, 2016), this microorganism was not found in the sprouts in the present study, neither in the study by Abadias et al. (2008), who evaluated the microbiological quality of minimally processed fruits and vegetables and sprouts from retail establishments in Spain. These results were different from those reported by Seo et al. (2010), who made a microbial evaluation in 112 samples of sprouts (broccoli, alfalfa, soybean, and clover) produced in Seoul, Korea, and found the presence of this microorganism in one sample.

It is worth noting that these bacteria are difficult to isolate since microorganisms capable of growing on the selective enrichment medium used for recovery of *L. monocytogenes* to the levels required for detection may be present in the food analyzed, preventing the growth of *L. monocytogenes* through simple competition. In addition, the levels of *L. monocytogenes* in sprouts may be lower than the threshold required for direct detection and recovery, thus resulting in false negative test results (Cauchon et al., 2017).

Chlorine sanitization treatments appear to be promising in reducing *L. monocytogenes* in seeds and sprouts. Iacumin and Comi (2019) inoculated *L. monocytogenes* strains on mung bean seeds and demonstrated that the levels of *L. monocytogenes* can be reduced after four washes in chlorinated water solution (100 ppm) from 3.6 log CFU/g to < 10 CFU in seeds, and from 9.8 log CFU/g to 1.9 log CFU/g in sprouts. In addition, Lee et al. (2002) investigated the inhibition by chemical treatment of *L. monocytogenes* inoculated into mung bean sprouts purchased from commercial stores in Washington and showed that the use of sodium hypochlorite (200 ppm) for 10 min was able to reduce the amount of the bacteria by 1.02 log CFU/g. In another work, treatment with aqueous chlorine dioxide (100 ppm) for 5 min reduced the population of *L. monocytogenes* inoculated into mung bean sprouts by 1.5 log CFU/g (Jin & Lee, 2007).

3.6 *Salmonella* spp.

In the present work, no *Salmonella* spp. strains were isolated in any of the analyzed samples, and this result is considered satisfactory according to the Brazilian legislation (Brasil, 2022)

and the European Commission regulation (European Commission, 2005), which determine that such strains must be absent. Similarly, Tornuk et al. (2011) investigated the presence of *Salmonella* spp. in wheat seed and sprouts, and Iacumin and Comi (2019) determined the quality of mung bean sprouts produced and sold in Italy between 2012 and 2016. Neither study found the presence of this microorganism.

However, in a trial aimed at evaluating the microbiological quality of 345 samples of minimally processed ready-to-eat vegetables, including 112 broccoli, alfalfa, soybean, and clover sprouts, Seo et al. (2010) found the presence of this microorganism in three samples of sprouts. In a survey by Saroj et al. (2006) that evaluated the microbiological quality of 124 sprouts marketed in India, *Salmonella* spp. was found in 24 samples. It is worth noting that, owing to the high risk of sprout-borne *Salmonella* infection, there is a need to improve the efficiency of detecting the presence of *Salmonella* spp. in sprouts, since this microorganism may be present at low concentrations or be sanitizer-injured (Zheng et al., 2015). Therefore, enrichment steps are considered as critical points in the detection of *Salmonella* spp. in sprouts, as they allow the rescue of cells of the bacterium to detectable levels (Zheng et al., 2015). Thus, in 2023, the FDA published an update of the Bacteriological Analytical Manual (Andrews et al., 2023), with new standards for the analysis of *Salmonella* spp. in sprouts, indicating the use of a specific broth for microbiological analysis of sprouts, the Universal Preenrichment Broth.

4 CONCLUSION

Disinfection with a household sanitizer reduced the microbial load of the sprout, especially as regards total coliforms and mesophilic aerobic bacteria. However, the microbial load remained high for both of them, which may have interfered with the possible detection of *L. monocytogenes* and *Salmonella* spp. Germination in the home environment seems to be a good strategy for the inclusion of a food item with high nutritional value in the diet; therefore, it is essential that seeds suitable for germination should be marketed.

In addition, national competent bodies need to create and disseminate a document of good production practices focused on the cultivation of sprouts, including in domestic environment, similar to the FDA—which is focused on industrial production—so that consumers can produce and safely consume this food at home from the microbiological point of view. Finally, further studies should be conducted to develop new sanitization techniques in the home environment that are effective in reducing the microbial load of sprouts as well as other fresh foods.

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