

Isolation, Identification and Partial Characterization of a *Lactobacillus casei* Strain with Bile Salt Hydrolase Activity from Pulque

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Abstract The aim of this study was to isolate, from pulque, *Lactobacillus* spp. capable of survival in simulated gastrointestinal stress conditions. Nine Gram-positive rods were isolated; however, only one strain (J57) shared identity with *Lactobacillus* and was registered as *Lactobacillus casei* J57 (GenBank accession: JN182264). The other strains were identified as *Bacillus* spp. The most significant observation during the test of tolerance to simulated gastrointestinal conditions (acidity, gastric juice and bile salts) was that *L. casei* J57 showed a rapid decrease ($p \leq 0.05$) in the viable population at 0 h. Bile salts were the stress condition that most affected its survival, from which deoxycholic acid and the mix of bile salts (oxgall) were the most toxic. *L. casei* J57 showed bile salt hydrolase activity over primary and secondary bile salts as follows: 44.91, 671.72, 45.27 and 61.57 U/mg to glycocholate, taurocholate, glycodeoxycholate and taurodeoxycholate. In contrast, the control strain (*L. casei* Shirota) only showed activity over tauroconjugates. These results suggest that *L. casei* J57 shows potential for probiotic applications.

Keywords Pulque · *Lactobacillus* · Bile salt hydrolase · Probiotics

Introduction

A fermented beverage known as “pulque” is produced by the fermentation of agave sap using its indigenous microorganisms [1, 2]. The agave sap is extracted from several species of maguey cactus such as *Agave atrovensis* and *A. Americana* [1, 2]. Slightly fermented agave sap is greatly appreciated in different regions of Mexico because it is popular belief that it provides health benefits in children, pregnant women and the elderly [1]. The preparation of this beverage involves three types of fermentation: acid, alcoholic and viscous. Therefore, the diversity of associated microorganisms is complex and a mixed culture of bacteria and yeasts such as *Zymomonas mobilis* [3], *Lactobacillus* spp., *Saccharomyces cerevisiae*, *Leuconostoc mesenteroides*, *Leuconostoc kimchii* [2], *Candida lusitanae* [4], *Kluyveromyces marxianus* [5], *Microbacterium arborescens*, *Flavobacterium johnsoniae*, *Acetobacter pomorium*, *Gluconobacter oxydans* and *Hafnia alvei* [6]. These microorganisms have been studied using molecular techniques [6], and it is known that some of them show potential biotechnological applications, such as those including the production of different compounds including lactic acid or ethanol, hydrolytic enzymes, and exopolysaccharides, and they also show applications in the health field [3, 6, 7]. In particular, the population of lactic acid bacteria (LAB) in pulque ranged from 6×10^7 to 2×10^{11} CFU/mL [1]. LAB have been identified as the main bacteria present in the early hours of fermentation of this beverage. LAB are a group of Gram-positive facultatively anaerobic bacteria, which excrete lactic acid as the

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main product of fermentation [6]. They are important microorganisms in some parts of the human and animal body (e.g., intestine, nasopharyngeal and vaginal mucosa) and also are found in milk, plants and fermented food such as sourdough, sauerkraut, boza [8] and pulque [2] among others. Such food and beverages have been and still are an important vehicle of living bacteria sources for the human body. Indeed, LAB strains from animal and human intestinal microbiota have been adopted as “probiotic” food supplements [9]. *Lactobacillus* is responsible for most of bile salt hydrolase activity detected in intestinal contents [10]. Bile salt hydrolases (BSHs) (EC 3.5.1.24) catalyze the hydrolysis of conjugated bile salts in the amide bond on the C-24 position of the steroid, which results in the production of a free amino acid (taurine or glycine) and an unconjugated bile acid molecule [11]. In recent years, the ability of some *Lactobacillus* to produce BSH has become the focus of attention on account of its influence on cholesterol metabolism, and hence, BSH activity can be explored as a functional probiotic biomarker for the selection of probiotic adjunct to manage hypercholesterolemia [10].

In spite of the fact that *Lactobacillus* in pulque has been identified by 16S rDNA analysis, it has not been isolated from this beverage and the behavior of these bacteria under gastrointestinal stress conditions where they face low pH; the presence of gastric juices and bile salts is still unknown. In order to offer health benefits, bacteria with probiotic potential have to be resistant to these stress conditions to reach the intestinal tract alive. The aim of the present study was to isolate strains of Lactobacilli from fermented agave sap (pulque) using a stress-inducing procedure to assess their BSH activity.

Materials and Methods

Agave Sap and Fermentation

Samples of freshly collected agave sap and overnight fermented agave sap were obtained from three different Mexican regions: Huitzilac in the State of Morelos (19°02'N, 99°16'W) with an altitude of 2550 m, Jocotitlan in the State of Mexico (19°44'N, 99°45'W) with an altitude of 3900 m and Singuilucan in the State of Hidalgo (19°59'N, 98°57'W) with an altitude of 2525 m. The samples were placed in sterile tubes and transported to the laboratory under a refrigerated condition of 4 °C. The fermentation of each sample was carried out in a 250-mL sterile glass container by addition of fermented agave sap and fresh agave sap at a ratio of 1:3 v/v (pulque) at 36 °C. The pH was determined after 3 h of fermentation.

Isolation Criteria, Bacterial Strains, Microbiological Media and Growth Conditions

In order to isolate bacteria capable of survival in simulated gastrointestinal stress conditions, 1 mL of pulque was added to a stress-inducing medium which consisted of Lactobacilli MRS Broth (pH 1.5, HCl 0.1 N) (Difco™, USA) and oxgall (0.5 % w/v) (Difco™, USA) and incubated for 2 h in a rotatory chamber (200 rpm, 37 °C) (Gallenkamp, UK). Afterward, 10 µL of the medium was plated onto Lactobacilli MRS Agar (Difco™, USA) and incubated for 12 h in an aerobic atmosphere at 37 °C. The resulting isolates and *L. casei* Shirota were observed under a light microscope. Rod-shaped microorganisms (bacilli) that were Gram positive were selected. In order to prepare enough experimental stock, the isolated microorganisms were plated onto Lactobacilli MRS Agar and incubated at 37 °C for 24 h; then, the biomass was transferred to Lactobacilli MRS Broth and incubated at 37 °C for 24 h. The broth containing the biomass was centrifuged at 30,000×g for 30 min at 4 °C under aseptic conditions. The cell pellet was suspended in 10 mL of Lactobacilli MRS Broth with 50 % glycerol and stored at –20 °C until use.

Pure culture of the probiotic bacteria *Lactobacillus casei* Shirota was isolated from Yakult® (Yakult Mexico), through growing it on Lactobacilli MRS Agar and incubating it for 24 h at 37 °C. In order to prepare enough experimental stock, biomass was transferred to Lactobacilli MRS Broth and incubated using the same conditions. Subsequently, the broth containing the biomass was centrifuged at 30,000×g for 30 min at 4 °C under sterile conditions. The cell pellet was suspended in 10 mL of Lactobacilli MRS Broth with 50 % glycerol and stored at –20 °C until use. *Lactobacillus casei* Shirota was used as a control for biochemical characteristics, antibiotic resistance, tolerance to simulated gastric juice and bile salts, and 16S rRNA sequencing [11]. This microorganism was grown in Mac Conkey Agar and incubated for 24 h at 37 °C. In order to prepare enough experimental stock, biomass was transferred to Mac Conkey Broth and the procedure mentioned above was followed.

Catalase Test

It was carried out in triplicate by placing a drop of hydrogen peroxide (J.T. Baker, USA) over a colony of microorganisms placed over a microscopic slide. If bubbles or froth were formed, the test was considered as positive [12]. *E. coli* ATCC 160211 was used as a positive control and *L. casei* Shirota as negative.

Hemolysis Test

The blood agar screening method of Ruiz-Moyano et al. [13], with slight modifications, was used. In brief, the strains were incubated overnight on Lactobacilli MRS Agar before the test. One colony of each strain was plated onto blood agar and incubated for 48 h at 37 °C in a brooder stove (Felisa[®], México). For each strain, the test was carried out in triplicate. *E. coli* ATCC 160211 was used as a positive control and *L. casei* Shirota as negative.

Carbohydrate Fermentation Pattern

The fermentation profile of isolated bacteria and *L. casei* Shirota was characterized by API 50 CHL test (bioMérieux, Marcy L'Etoile, France) [14]. The fermentation profile obtained was evaluated with *apiweb*[™] software. *Lactobacillus casei* Shirota was used as positive control and *E. coli* ATCC 160211 a negative.

Genotypic Identification

DNA extraction was performed using Easy-DNA[™] kit for genomic DNA isolation (Invitrogen, Cergy-Pontoise, France) according to the supplier's instructions. PCR amplification of the isolated strains was carried out using primers LAC1 (5'-AGCAGTAGGGAATCTTCCA-3') and LAC2 (5'-ATTCACCGCTACACATG-3') derived for the amplification of 340 bp of the 16S rDNA gene of *Lactobacillus* [15]. Amplifications were performed with a thermal cycler T gradient (Promega) using the conditions reported by Walter et al. [15]. PCR-amplified products were purified with QIAquick PCR purification kit 50 (QIAGEN, USA) and analyzed by 2 % agarose gel electrophoresis, ethidium bromide staining under UV light. DNA sequences were determined by using the genetic analyzer ABI PRISM[®] 3100 (Applied Biosystems, USA). Comparisons and sequence alignments were made by using BioEdit DNA sequences analyzer (<http://www.mbio.ncsu.edu/BioEdit/BioEdit.html>) [16] and the basic local alignment search tool (<http://blast.ncbi.nlm.nih.gov/Blast.cgi>).

Tolerance to Acid, Simulated Gastric Juice and Bile Salts

Simulated gastrointestinal stress conditions were acidity, gastric juice and bile salts. The tolerance of the isolated microorganism to these conditions was measured at different times (0, 1, 2 and 4 h). Acid tolerance was studied by inoculating an aliquot of approximately 10⁸ CFU/mL into sterile acid water [17]. This water was adjusted with HCl (0.1 M) (J.T. Baker, USA) to 1.5 pH by using a pH meter (Orion 410A+, Thermo Scientific, USA). Gastric

juice tolerance was determined according to Vinderola and Reinheimer [18] with slight modifications, using a solution containing pepsin (0.3 % w/v) (SIGMA, USA) and NaCl (0.5 % w/v) (J.T. Baker, USA) adjusted to pH 2.0 using HCl (0.1 M) (J.T. Baker, Mexico). Bile salt tolerance was tested by using bovine bile (oxgall, Difco[™]), primary bile acids conjugated either taurine (taurocholic, TC) or glycine (glycocholic acid, GC) (SIGMA, USA) and a secondary bile acid (deoxycholic acid, DC) (SIGMA, USA). Bile salt tolerance of each salt was studied using Lactobacilli MRS Broth supplemented with 0.3 % w/v of bile salt [6]. In all tolerance tests, decimal dilutions in peptone and water solution (0.5 % w/v) were individually stirred for 30 s. Afterward, at each time of exposure, a sample of 100 µL was taken, plated onto Lactobacilli MRS Agar and incubated in an aerobic atmosphere for 48 h at 37 °C. Tolerance was determined by comparing the plate counts at different times with the initial one at time zero. The initial count (approximately 10⁸ CFU/mL) with no treatment was considered 100 % survival. Zero time (0 h) with treatment only means the time in which bacteria were poured to the stressful condition and the sample was taken. The results were normalized by using Eq. (1) and expressed relative survival of strains on Lactobacilli MRS Agar. All of the experiments were carried out in triplicate.

$$\%RS = \left(\frac{\log \text{CFU after treatment}}{\log \text{CFU initial count}} \right) \times 100 \quad (1)$$

Antibiotic Resistance Test

BIO-RAD multidisc (Gram-positive II bacteria) containing cephalothin (30 µg), cefotaxime (30 µg), levofloxacin (5 µg), cefuroxime (30 µg), dicloxacillin (1 µg), erythromycin (15 µg), gentamicin (10 µg), cefepime (30 µg), penicillin (30 µg), tetracycline (30 µg), ampicillin (10 µg) and trimethoprim-sulfamethoxazole (25 µg) was employed for antibiotic resistance tests. The multidisc was placed onto Muller Hinton agar plates previously inoculated with *L. casei* J57 or *L. casei* Shirota (Yakult[®]). Plates were incubated for 72 h at 37 °C in an anaerobic chamber (Forma anaerobic system Model 1025, Thermo Scientific, USA) with an atmosphere of 10 % CO₂, 5 % H₂ and 85 % N₂ [19]. The test was carried out in triplicate, and the results were expressed according to the instructions given by the supplier of the test kit. Before the test, we used *E. coli* ATCC 160211 to measure the reproducibility of the technique (data not shown).

Bile Salt Hydrolase Test

Bile salt hydrolase (BSH) assay was carried out in spent broth as described by González-Vázquez et al. [11] using

overnight cultures of *L. casei* J57 or *L. casei* Shirota (Yakult®). The activity of each strain was evaluated in triplicate. One unit of BSH activity was defined as the amount of enzyme which liberated 1 mmol of amino acids per 1 mL of substrate per minute per mg of protein, determined by the Bradford method (Bio-Rad, Mexico) and according to the supplier's recommendations. Each assay was carried out in triplicate.

Results

Isolation and Identification (Phenotypic and Genotypic) of Bacteria from Agave Sap

The pH after 3-h fermentation was 4.6 ± 0.06 , 4.15 ± 0.05 and 3.7 ± 0.06 in the pulque of Singuilucan, Jocotitlán and Huitzilac regions, respectively. The stress-inducing medium allowed us to isolate 76 microorganisms from pulque of the three Mexican regions tested. However, only nine isolates were Gram-positive rod-shaped bacteria. Therefore, they were chosen for further investigation. Four of these nine bacteria were isolated from Singuilucan (S67, S37, S38 and S51), one from Jocotitlán (J57) and four from Huitzilac (H18, H19, H64 and H60). The strain J57 was catalase negative and non-hemolytic, and the other eight strains were catalase positive and hemolytic. The API 50 CHL test determined that *L. casei* Shirota and *L. casei* J57 had the ability to ferment the sugars galactose, lactose and sorbose and the sugar alcohol sorbitol. Other sugars that were fermented by both organisms were D-ribose, D-glucose, D-fructose, D-mannose, D-mannitol, D-sorbitol, D-cellobiose, D-maltose, D-lactose, D-trehalose, genentiobiose, D-turanose, D-tagatose, N-acetyl glucosamine, amygdalin, arbutin, esculin ferric citrate, salicin and potassium gluconate. They only differed in the fermentation of D-adonitol. Therefore, the strain J57 was identified as *L. casei* subsp. *paracasei* (99.7 %). The other isolated strains did not exhibit identity with the genus reported in the API database, since this system is designed to identify bacteria only belonging to the *Lactobacillus* genus. Therefore, these last strains were discarded for the rest of the tests. The BLAST analysis was used for genotypic identification. The 16S rRNA sequence of the strain shared 99 % identity with that of *L. casei* (GenBank accession: JN182264).

Antibiotic Resistance

In the antibiotic tests, the strain *L. casei* J57 showed resistance to cefepime and sensitive to the other antibiotics tested. In contrast, *L. casei* Shirota showed resistance to dicloxacillin and gentamicin and sensitive to the other antibiotics tested.

Tolerances

Acidity

Regarding acids, simulated gastric juice and bile salt tolerance, the most significant observation was a significant decrease ($p \leq 0.05$) in a viable count from the initial population (0 h). The relative survival percentage of *L. casei* J57 and the *L. casei* Shirota at pH 1.5 over 4 h is shown in Fig. 1a. Under such conditions, *L. casei* J57 was able to survive for over 4 h. In contrast, the *L. casei* Shirota survived for 1 h.

Simulated Gastric Juice

The relative survival percentage under gastric juice of *L. casei* J57 and *L. casei* Shirota is shown in Fig. 1b. *L. casei* J57 survived for 4 h under the experimental conditions. Conversely, *L. casei* Shirota did not show viability after 1 h.

Bile Salts

Lactobacillus casei J57 and the *L. casei* Shirota survived for 4 h of exposure to TC, GC, DC and oxgall (Fig. 1c–f, respectively). It should be noted that oxgall had a higher negative effect on the percentage of relative survival than the other bile salts tested.

Bile Salt Hydrolase Activity

Figure 2 shows the BSH activity of *L. casei* J57 using different bile salts (GC: 44.91; GDC: 45.27; TC: 671.72; TDC: 61.57 U/mg of protein) and the BSH activity of *L. casei* Shirota, which did not show any activity in glycine conjugates. However, its activity over tauroconjugates (TC: 1046.15; TDC: 264.69 U/mg of protein) was higher than the ones showed by *L. casei* J57. In both cases, hydrolase activity toward taurocholic acid was higher than the activity shown to other bile salts.

Discussion

In order to determine whether a bacterium has the ability to survive under gastrointestinal tract conditions, it is important to evaluate its ability to survive under such stress [9]. Therefore, the results obtained in the present work suggest that the use of a stress-inducing medium, which simulates similar conditions to those found in the gastrointestinal tract, allowed microorganisms that tolerate these conditions to be isolated. Particularly, this could be associated with the BSH activity shown by *L. casei* J57,

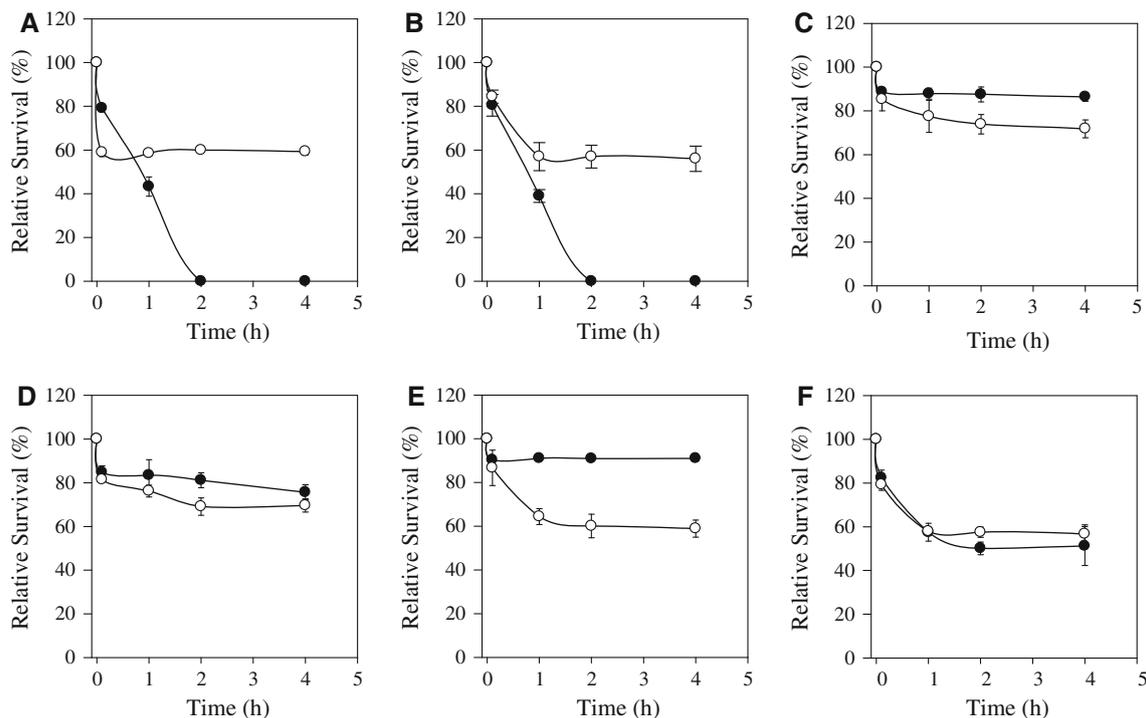


Fig. 1 Tolerances to the different simulated gastrointestinal stress conditions tested: **a** acidity, **b** gastric juice, **c** taurocholic acid, **d** glycocholic acid, **e** deoxycholic acid and **f** oxgall of *L. casei* J57

(circle) and *L. casei* Shirota (Yakult[®]) (filled circle). Values are the means of data generated from triplicate samples examined in three independent experiments

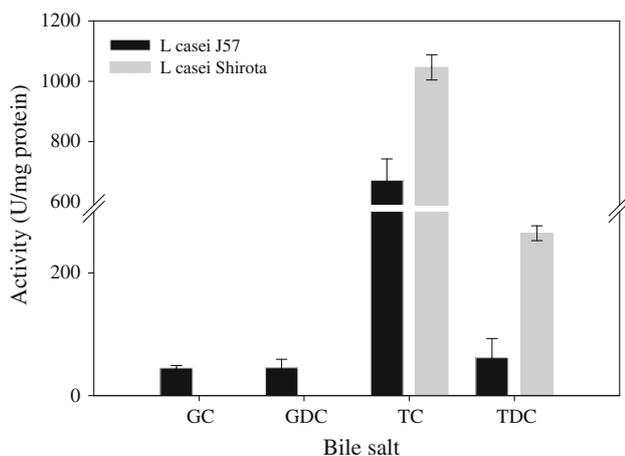


Fig. 2 Bile salt hydrolase activity in the spent broth of *L. casei* J57 and *L. casei* Shirota (Yakult[®]) grown in MRS broth supplemented with 0.5 % of glycocholic acid (GC), glycodeoxycholic acid (GDC), taurocholate (TC) and taurodeoxycholate (TDC). Values are the means of data generated from triplicate samples examined in three independent experiments

since it was isolated from a medium that contained bile salts, and cross-resistance mechanisms could be present.

Many LAB have been found in several fermented native foods [8]. In agave sap, some *Lactobacillus* have been identified as *L. acidophilus*, *L. kefir*, *L. acetotolerans*, *L.*

hilgardii, *L. plantarum*, *L. acidophilus*, *L. hilgardii*, *L. paracollinoides* and *Lactobacillus* spp. Y10c7 [6]. Particularly, *L. casei* is a ubiquitous microorganism, which is found in fermented dairy products, fresh vegetables and human sources [8]. Nevertheless, this is the first time that *L. casei* has been isolated and identified from pulque, and the first time that its BSH activity has been determined. The carbohydrate fermentation pattern between *L. casei* Shirota and *L. casei* J57 was similar as we expected; however, they differed in the fermentation of one carbohydrate, which indicate that probably they are different subspecies.

Regarding the tolerances tested, acidity is an environmental condition that is commonly found in the human gastrointestinal tract [20]; gastric pH increases during food intake from 1.5 to 3.0 or 5.0 [21]. In the case of beneficial bacteria, good acid tolerance is desirable, since it is usually related by cross-resistance to some other stress factor of the intestinal environment [22]. In this study, during the acid tolerance test, it was not possible to obtain colony-forming units of the control at the end of the experiment (Fig. 1a) because this condition (pH 1.5) was extremely stressful to this microorganism. Another important issue to consider is that after a meal, food ingredients can exert a protective effect over *Lactobacillus* strains [23]. In spite of these results, *L. casei* J57 (Fig. 1a) showed viability until the end

of the experiment, which indicates that this microorganism could survive this stress in the gastrointestinal tract.

Related to the tolerance to gastric juice, it was found that *L. casei* Shirota, which showed 30–40 % of tolerance to acidity, showed a similar behavior when it was under gastric juice; in contrast, *L. casei* J57 showed a higher tolerance to gastric juice (50–55 %) (Fig. 1b) and survived for 4 h under this stress. These results could indicate that a cross-resistance mechanism is present, since most pH stress protection systems in bacteria include a mechanism for sustaining cytoplasmic pH, and many pH stress-inducible systems offer cross-protection to other stresses such as increasing salt tolerances [22]. In addition, Todorov et al. [24] have suggested that resistance to low pH and elevated concentrations of bile salts is important for the growth and survival of bacteria in the intestinal tract.

Another important finding with regard to bile salts was that glycine conjugates have a stronger effect over strain survival (Fig. 1d) than taurine conjugates (Fig. 1c). De Smet et al. [25] reported that glycine conjugates are far more toxic than taurine conjugates due to their different pKa (3.9 and 1.0, respectively). On the other hand, oxgall had a larger effect than the previously tested salts over the viability of *L. casei* J57 and *L. casei* Shirota (Fig. 1f), suggesting a toxic synergistic effect when the strains are in contact with the mixture of bile salts, since oxgall is manufactured from large quantities of fresh bile by the rapid evaporation of the water content and is made up of fatty acids, bile acids, inorganic salts, sulfates, bile pigments, cholesterol, mucin, lecithin, glucuronic acids, porphyrins and urea [26, 27].

In the present work, the antibiotic resistance profile of *L. casei* J57 matched the profile reported for other *Lactobacillus* [28]. However, it showed resistance to cefepime (a fourth-generation cephalosporin). We did not find any previous studies that tested cefepime susceptibility [29], and most of the studies have included only a limited number of *Lactobacillus* strains or antibiotics. This is probably because *Lactobacillus* has been considered as generally recognized as safe (GRAS), and therefore, their antimicrobial susceptibility has received only little attention [29]. Further studies are needed in support of the resistance of *Lactobacillus* to fourth-generation cephalosporin. At this stage, cefepime is contraindicated in patients with known allergies to cephalosporin and or penicillin antibiotic and is usually reserved to treat moderate to severe nosocomial pneumonia infections [30]. Additionally, the strain J57 was identified as *Lactobacillus* with non-hemolytic activity, which is particularly important since the presence of this activity can contribute to the virulence of some pathogenic bacteria [31, 32]. However, this effect depends on different factors, such as whether the host is immunocompromised [33].

Regarding BSH activity (Fig. 2), it was only determined in the case of strain *L. casei* J57 since this activity has been related to *Lactobacillus* species with a possible probiotic application, due to the association of BSH activity with serum cholesterol-lowering effects [9]. *L. casei* J57 showed BSH activity over primary and secondary bile salts conjugated to glycine or taurine (Fig. 1). However, lower activity was shown than for *L. casei* (control) regarding either primary or secondary tauroconjugates. In spite of the activity found by a quantitative method, some authors [34] have reported the lack of BSH activity by using a plate assay over tauroconjugates by several *L. casei* strains, including the one isolated from a popular probiotic drink (Yakult®). Conjugated bile salts are periodically released into the intestinal environment. According to Begley et al. [35], the ratio of glycine conjugates to tauroconjugates in human bile is usually 3:1, which shows the importance of BSH activity over glycine conjugates from *Lactobacillus* with probiotic features. It is known that bile salts are toxic to bacteria [25]. Therefore, bacteria in the intestine may express BSH activity to protect them from such toxicity. These results may explain why *L. casei* Shirota in the present work showed tolerance to different glycine conjugated bile salts but did not show hydrolase activity toward such salts.

We conclude that the use of a stress-inducing procedure allows the isolation of lactic acid bacterial strains capable of showing tolerance to gastrointestinal stress and BSH activity, which were higher than those of the commercial probiotic used in our study. This activity is considered as a probiotic characteristic, which could impact on health.

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Compliance with Ethical Standards

Conflict of interest The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

References

1. Valadez-Blanco R, Bravo-Villa G, Santos-Sánchez N, Velasco-Almendarez S, Montville T (2012) The artisanal production of pulque, a traditional beverage of the Mexican highlands. *Probiotics Antimicrob Proteins* 4(2):140–144. doi:10.1007/s12602-012-9096-9
2. Escalante A, Giles-Gomez M, Hernandez G, Cordova-Aguilar MS, Lopez-Munguia A, Gosset G, Bolivar F (2008) Analysis of bacterial community during the fermentation of pulque, a traditional Mexican alcoholic beverage, using a polyphasic approach. *Int J Food Microbiol* 124(2):126–134. doi:10.1016/j.ijfoodmicro.2008.03.003
3. Correa-Ascencio M, Robertson IG, Cabrera-Cortés O, Cabrera-Castro R, Evershed RP (2014) Pulque production from fermented

- agave sap as a dietary supplement in Prehispanic Mesoamerica. *Proc Natl Acad Sci* 111(39):14223–14228
4. Lira A, Alvarado-Resendiz M, Simental S, Martini J, Reyes-Santamaria M, Guemes-Vera N (2014) Use of *Lactobacillus* from pulque in sourdough. *Adv Microbiol* 4:969–977. doi:10.4236/aim.2014.414108
 5. Estrada-Godina AR, Cruz-Guerrero AE, Lappe P, Ulloa M, García-Garibay M, Gómez-Ruiz L (2001) Isolation and identification of killer yeasts from agave sap (aguamiel) and pulque. *World J Microbiol Biotechnol* 17(6):557–560. doi:10.1023/A:1012210106203
 6. Escalante A, Rodriguez ME, Martinez A, Lopez-Munguia A, Bolivar F, Gosset G (2004) Characterization of bacterial diversity in pulque, a traditional Mexican alcoholic fermented beverage, as determined by 16S rDNA analysis. *FEMS Microbiol Lett* 235(2):273–279. doi:10.1016/j.femsle.2004.04.045
 7. Chellapandian M, Larios C, Sanchez-Gonzalez M, Lopez-Munguia A (1998) Production and properties of a dextranucrase from *Leuconostoc mesenteroides* IBT-PQ isolated from ‘pulque’, a traditional Aztec alcoholic beverage. *J Ind Microbiol Biotechnol* 21(1–2):51–56. doi:10.1038/sj.jim.2900560
 8. Rivera-Espinoza Y, Gallardo-Navarro Y (2010) Non-dairy probiotic products. *Food Microbiol* 27:1–11
 9. Patel AK, Singhania RR, Pandey A, Chincholkar SB (2009) Probiotic bile salt hydrolase: current developments and perspectives. *Appl Biochem Biotechnol* 162(1):166–180. doi:10.1007/s12010-009-8738-1
 10. Kumar R, Grover S, Batish V (2012) Bile salt hydrolase (Bsh) activity screening of lactobacilli: in vitro selection of indigenous lactobacillus strains with potential bile salt hydrolysing and cholesterol-lowering ability. *Probiotics Antimicrob Proteins* 4(3):162–172. doi:10.1007/s12602-012-9101-3
 11. González-Vázquez R, Gutiérrez-López GF, Arellano-Cárdenas S, López-Villegas EO, Téllez-Medina DI, Rivera-Espinoza Y (2014) Morphometric parameters, zeta potential and growth rate of *Lactobacillus casei* Shirota by effect of different bile salts. *Rev Mex Ing Quim* 13:189–199
 12. MacFaddin JF (1980) *Biochemical tests for identification of medical bacteria*. Lippincott Williams and Wilkins, Philadelphia
 13. Ruiz-Moyano S, Martin A, Benito MJ, Casquete R, Serradilla MJ, Cordoba Mde G (2009) Safety and functional aspects of pre-selected lactobacilli for probiotic use in Iberian dry-fermented sausages. *Meat Sci* 83(3):460–467. doi:10.1016/j.meatsci.2009.06.027
 14. Annuk H, Shchepetova J, Kullisaar T, Songisepp E, Zilmer M, Mikelsaar M (2003) Characterization of intestinal lactobacilli as putative probiotic candidates. *J Appl Microbiol* 94(3):403–412
 15. Walter J, Hertel C, Tannock GW, Lis CM, Munro K, Hammes WP (2001) Detection of *Lactobacillus*, *Pediococcus*, *Leuconostoc*, and *Weissella* species in human feces by using group-specific PCR primers and denaturing gradient gel electrophoresis. *Appl Environ Microbiol* 67(6):2578–2585. doi:10.1128/aem.67.6.2578-2585.2001
 16. Thomas AH (1999) BioEdit: a user-friendly biological sequence alignment editor and analysis program for Windows 95/98/NT. *Nucleic Acids Symp Ser* 41:95–98
 17. Mishra V, Prasad DN (2005) Application of in vitro methods for selection of *Lactobacillus casei* strains as potential probiotics. *Int J Food Microbiol* 103(1):109–115. doi:10.1016/j.ijfoodmicro.2004.10.047
 18. Vinderola CG, Reinheimer JA (2003) Lactic acid starter and probiotic bacteria: a comparative “in vitro” study of probiotic characteristics and biological barrier resistance. *Food Res Int* 36(9–10):895–904. doi:10.1016/S0963-9969(03)00098-X
 19. Melgar-Lalanne G, Rivera-Espinoza Y, Farrera-Rebollo R, Hernández-Sánchez H (2014) Survival under stress of halotolerant lactobacilli with probiotic properties. *Rev Mex Ing Quim* 13:323–335
 20. Mayorga-Reyes L, Bustamante-Camilo P, Gutiérrez-Nava A, Barranco-Florido E, Azaola-Espinosa A (2009) Crecimiento, sobrevivencia y adaptación de *Bifidobacterium infantis* a condiciones ácidas. *Rev Mex Ing Quim* 8:259–264
 21. Cotter PD, Hill C (2003) Surviving the acid test: responses of gram-positive bacteria to low pH. *Microbiol Mol Biol Rev* 67(3):429–453
 22. Beales N (2004) Adaptation of microorganisms to cold temperatures, weak acid preservatives, low pH, and osmotic stress: a review. *Compr Rev Food Sci Food Saf* 3(1):1–20. doi:10.1111/j.1541-4337.2004.tb00057.x
 23. Morelli L (2000) In vitro selection of probiotic lactobacilli: a critical appraisal. *Curr Issues Intest Microbiol* 1(2):59–67
 24. Todorov SD, Botes M, Guigas C, Schillinger U, Wiid I, Wachsmann MB, Holzapfel WH, Dicks LM (2008) Boza, a natural source of probiotic lactic acid bacteria. *J Appl Microbiol* 104(2):465–477. doi:10.1111/j.1365-2672.2007.03558.x
 25. De Smet I, Van Hoorde L, Vande Woestyne M, Christiaens H, Verstraete W (1995) Significance of bile salt hydrolytic activities of lactobacilli. *J Appl Bacteriol* 79(3):292–301
 26. Isenberg HD (1992) *Clinical microbiology procedures handbook*, vol 1. American Society for Microbiology, Washington DC
 27. Murray PR, Baron EJ, Pfaller MA, Tenover FC, Tenover RH (1995) *Manual of clinical microbiology*. American Society for Microbiology, Washington DC
 28. Mourad K, Nour-Eddine K (2006) In vitro preselection criteria for probiotic *Lactobacillus plantarum* strains of fermented olives origin. *Int J Probiotics Prebiotics* 1:27–32
 29. Salminen MK, Rautelin H, Tynkkynen S, Poussa T, Saxelin M, Valtonen V, Jarvinen A (2006) *Lactobacillus* bacteremia, species identification, and antimicrobial susceptibility of 85 blood isolates. *Clin Infect Dis* 42(5):e35–e44. doi:10.1086/500214
 30. Yahav D, Paul M, Fraser A, Sarid N, Leibovici L (2007) Efficacy and safety of cefepime: a systematic review and meta-analysis. *Lancet Infect Dis* 7(5):338–348. doi:10.1016/s1473-3099(07)70109-3
 31. Chang CI, Liu WY, Shyu CZ (2000) Use of prawn blood agar hemolysis to screen for bacteria pathogenic to cultured tiger prawns *Penaeus monodon*. *Dis Aquat Organ* 43(2):153–157. doi:10.3354/dao043153
 32. Sritharan M (2006) Iron and bacterial virulence. *Indian J Med Microbiol* 24(3):163–164
 33. Borriello SP, Hammes WP, Holzapfel W, Marteau P, Schrezenmeier J, Vaara M, Valtonen V (2003) Safety of probiotics that contain lactobacilli or bifidobacteria. *Clin Infect Dis* 36(6):775–780. doi:10.1086/368080
 34. Dashkevich MP, Feighner SD (1989) Development of a differential medium for bile salt hydrolase-active *Lactobacillus* spp. *Appl Environ Microbiol* 55(1):11–16
 35. Begley M, Gahan CG, Hill C (2005) The interaction between bacteria and bile. *FEMS Microbiol Rev* 29(4):625–651. doi:10.1016/j.femsre.2004.09.003