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Article

Shelf-Life Study of Probiotic Dahi Prepared by a New Isolated Probiotic, Lactobacillus acidophilus Strain JB2CON

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Abstract: The principal goal of this current study was to isolate a new potential probiotic strain to prepare contamination-free probiotic dahi in a dosage form with desired shelf life. To achieve our target, firstly, probiotic dahi was prepared by a probiotic strain, *Lactobacillus acidophilus* JB2CON (OM909067) (identified by 16S sequencing) isolated from natural dahi in Dhaka, Bangladesh. The isolate showed five probiotic features by *in vitro* test and had an antagonistic effect against five different pathogens. More interestingly, it inhibited *Escherichia coli* (diameter of zone of inhibition = 15.75±0.35) the most. In the time-kill analysis method, *E. coli* was killed totally within 5.50 hours of fermentation. Finally, the time duration of 14 days was an ideal shelf life, because the probiotic count increased but a slight change in pH did not affect the taste (sourness/acidity) of dahi. The most significant impact of this research is that the probiotic dahi of our experiment might be used as a cheap option for preventing pathogenic *Escherichia coli*-related foodborne diseases worldwide, especially in underdeveloped countries like Bangladesh.

Keywords: L. acidophilus JB2CON; probiotic dahi; shelf life; Escherichia coli

1. Introduction

Probiotics are beneficial for the health of humans. The most popularly consumed probiotics are Lactobacillus acidophilus (Anjam et al., 2014). Research has demonstrated that the presence of L. acidophilus can produce some potential probiotic effects in humans, such as; performing as a barrier against pathogens, recovering lactose intolerance, boosting immune response, and decreasing cholesterol levels (Alberts, 1987). L. acidophilus can colonize the intestines of humans and inhibit pathogens, such as Escherichia coli, Salmonella typhimurium, Streptococcus aureus, and so on. These pathogens also inhabit the intestines (Anjam et al., 2014, Rolhion at al., 2015). These probiotics are applied to prevent diarrheal infections as well (Britton et al. 2009). According to the last Global Burden of Disease Study, approximately 2.39 billion people were infected by diarrheal cases globally and nearly 0.53 million under-five children died yearly. In lower and middle-income (LMI) countries, specifically, Bangladesh, incidence and case-fatality ratios are much higher as compared with that of developing countries (Sarker et al., 2018). In Bangladesh, the average cost per episode was US \$ 67.18, while the moderate inpatient and outpatient charge were US \$ 110.51 and US \$ 23.62 respectively. The cost was substantially largest for impoverished households, 21.45% of household income, compared to 4.21% of the prosperous people (Sarker et al., 2018). L. acidophilus can be a cost-saving option for preventing diarrheal disease and pathogenic Escherichia coli-related other food-borne diseases of billions of people around the world.



Potential probiotic strains are vital for killing pathogens in the intestines. In Bangladesh most of all, marketed probiotics are purchased from foreign countries. There is very little research on isolated probiotics in Bangladesh (Anjuman et al., 2015). Moreover, assessment of the potential probiotic traits of a strain is essential for claiming it as a probiotic strain (Sarker et al., 2018; Chidre et al., 2017). Isolation of potential probiotics from Bangladeshi natural sources is significant to boost the immunity of the masses of this geographic territory. In this study, we focused on the isolation of a new probiotic strain of *L. acidophilus* from Bangladeshi natural sources.

Probiotic dahi is a good option for delivering probiotics to the intestines. Dahi is a customary <u>yogurt</u> or <u>fermented milk product</u> in the <u>Indian subcontinent</u>, usually produced from <u>cow's milk</u>, and also from <u>buffalo milk</u>, or <u>goat milk</u>. It is consumed all around the Indian subcontinent (Caballero et al., 2015). We can get two benefits from probiotic dahi: one is a test, and another one is probiotic culture. In the Bangladeshi market, there are no probiotic dahi having levels of specific probiotic bacteria (Anjuman et al., 2015).

The shelf life of dahi is another important factor for maintaining the quality of dahi up to the expiration date. There is a correlation between pH, temperature, probiotic growth, and contamination in dahi (Schmidt et al., 1996; Yang et al., 2018). The preparation of dahi in unhygienic environments and contaminated starter cultures can spoil dahi and cause food poison as well. Shelf life is dependent on this issue. Shelf life depends on another factor, that is, total live probiotic count Guesh et al., 2020; Fijan, 2016).

The main target of this topical investigation was to prepare a probiotic dahi having potential probiotic *L. acidophilus* in dosage form, that is, the number of probiotics would not be less than a certain number of colony forming units per gram of dahi. Secondly, we aimed to find the capability of dahi for inhibiting the growth of contaminations. Information on probiotic growth and inhibition of microbial contamination would be used for determining shelf life of dahi.

2. Materials and Methods

2.1. Probiotic isolation and identification

2.1.1. Probiotic isolation

Probiotic species were isolated from a number of samples of Dahi in Dhaka, Bangladesh. 1 mL of 10-9 decimal diluted sample (suspended in normal 0.9% (w/v) saline solution) was spread on 20-25mL MRS (de Man Rogosa Sharpe, Oxoid, UK) agar medium, and the plates were incubated for 24-72 h in 37°C. Based on morphology, the distinguished most common bacterial colonies were selected from MRS agar media. To purify colonies, the isolates were streaked on the same media and finally, the pure colonies were transferred to MRS broth with 15% glycerol for further research. The Leica MZ9.5 (Germany), a potent stereo microscopic instrument that features a fantastic 9.5:1 zoom ratio and magnification capabilities as high as 480x, was utilized to analyze the colony morphology.

2.1.2. Gram staining test, 16S gene sequencing, and phylogenetic tree

The selected colonies were examined using gram staining protocol according to Coica (2005) method, and then observed under light microscope by 100X resolution (Coico, 2005). The MRS-agar cultured colonies were used for genomic DNA isolation with Phenol chloroform chemical lysis method according to the manufacturer's protocol. After DNA extraction, the concentration and purity of DNA was checked using Nanodrop® spectrophotometer ND2000 (Thermo Scientific, USA). The 260/280 ratio (absorbance at 260nm and 280nm) provided an indication of the purity of the DNA. The ratio of the sample obtained is above 1.8. In this study, 16S rDNA amplification and sequencing were performed based on the methodology described previously by Rahman et al. (2017). The universal 16S rRNA primer set for polymerase chain reaction (PCR) amplification was as follows: 27f (5'AGAGTTTGATCCTGGCTCAG-3'and 1492r (5'- GGTTACCTTGTTACGACTT-3') (Rahman et al., 2017).

The phylogenetic tree was inferred using the Neighbor-Joining method (Saitou and Nei, 1987). The bootstrap consensus tree inferred from 1000 replicates is taken to represent the evolutionary history of the taxa analyzed (Felsenstein, 1985). This analysis involved 12 nucleotide sequences. Evolutionary analyses were conducted in MEGA11 (Tamura et al., 2021).

2.1.3. Cellular morphology analysis under Transmission Electron Microscope

With a plasma cleaner (Power: Low, 12 sec), we hydrophilized the grid beforehand. After that, a tabletop centrifuge was put to work to spin the bacterial culture for two minutes at 2,000–4,000 rpm. Ten microliters of pure water were used for the re-suspension of the particle. After applying the specimen to the grid, it was left in place for one minute. The samples had been incubated for one minute, stained with a 1.8% uranyl acetate solution, and then allowed to air dry. The photos of bacterial samples were captured with a JEM 1010 electron transmission microscope (JEOL).

2.2. Probiotic potentiality and safety test

2.2.1. Antibiotic sensitivity test by agar well diffusion method

An antibiotic sensitivity test was conducted against the probiotic isolate JB2CON following the method of Chetan et al. (2017) with some modifications. 8 mL MRS agar (with log 8 CFU/mL of the isolate JB2CON) overplayed on previously solidified MRS agar (Chetan et al., 2017). Cefuroxim (100µl/30µg), Gentamycin (100µl/10µg), Chloramphenicol (100µl/30µg), Erythromycin (15 µg / 100 µL), Clindamycin (10 µg / 100 µL), Tylosin (30 µg / 100 µL), Amphicilin (10 µg / 100 µL), Vancomycin (30 µg / 100 µL), Kanamycin (30 µg / 100 µL), Streptomycin (10 µg / 100 µL), and Tetracycine (30 µg / 100 µL) were filled in a 7mm diameter well in MRS agar media with an upper layer of the probiotic bacterial strain. The test Petri dishes were incubated for 24 hours at 37°C. Antibiotic sensitivity for every antibiotic was tested twice.

2.2.2. Hemolysis test

To evaluate for hemolytic activity, blood agar (BD, USA) plates were streaked with MRS broth that contained JB2CON strain cultures and underwent incubation for 72 hours at 37 °C. Next, each of the plates were inspected to see if the *Lactobacillus* colonies were surrounded by any greenish (α -hemolysis) or clean (β -hemolysis) hemolytic zones, or if there was none at all (γ -hemolysis). The test was conducted for three times.

2.2.3. Acid tolerance

The acid tolerance of the isolate JB2CON was tested following the method of Ortakci et al. (2012) with some modifications (Ortakci et al., 2012). 0.2g NaCl plus 0.7 mL HCl (Stock solution), and sufficient water were used to make 100 mL simulated gastric juice (SGJ of pH 1.4). SGJ and sterilized dahi were mixed in a ratio of 1:4 for making a solution of pH 2.35 (pH <3) (Ortakci et al., 2012) This final solution was used for analyzing the capability of the isolate JB2CON *in* our study to tolerate acidity (pH <3). 1 mL probiotic isolate (1.28×10⁸ CFU/mL) was added to 9 mL final solution of SGJ and dahi. Average CFU of the probiotic from two tests was counted after 24 hours of incubation at 37°C.

2.2.4. Bile salt tolerance

With minor adjustments, the bile tolerance of the isolate JB2CON was assessed using the methodology of Hassanzadazar et al. (2012) (Hassanzadazar et al., 2012). The isolate JB2CON was incubated for 24 hours at 35°C in MRS broth. Following this time, the cells were suspended using a slow vortex. After that, 1% cell suspension broth was mixed with the MRS broth containing bile salt (0.5% and 0.25% ox-gall). The isolate JB2CON's viability was assessed after 8 hours of incubation by culturing it in petri-dishes using the spread plate technique, which was then incubated for 5 days at

35°C. The control was the cell suspension broth that had been incubated at 35°C for 8 hours without any bile salt added. Two replicates were used for each test.

2.2.5. Antimicrobial activity test by agar well diffusion method

Antimicrobial activity test of the isolate JB2CON was conducted following the method of Chidre et al. (2017) with some modifications. Antagonistic effects against six pathogenic microorganisms were tested in TSA media (Chidre et al., 2017). 100µl supernatant of the cell suspension broth of the isolate JB2CON (final pH 3.98 after 48 hours incubation at 35°C) was placed in a 7mm diameter well in TSA media having a lawn of pathogenic microorganism. Two replicates of antagonistic tests were conducted against each pathogen. Salmonella typhimurium ATCC 14028, Staphylococcus aureus ATCC6538, Escherichia coli ATCC 8739, Bacillus subtitlis ATCC 6633, Pseudomonas aeruginosa ATCC 142, and Candida albicans ATCC 10231 were bought from a local supplier in Dhaka, Bangladesh. Antimicrobial activity against these pathogens was tested twice.

2.2.6. Antimicrobial activity test by time-kill assay against unknown contamination

In the agar diffusion assay we used specific pathogens of ATCC cultures, although in the time-kill assay we employed unknown contaminations. Antimicrobial activity test was performed with some modifications by time-kill assay described by Prabhurajeshwar and Kelmani (2019) (Prabhurajeshwar and Kelmani, 2019). For this assay, random contaminations and all ingredients for dahi (pasteurized milk, sugar, plus the isolate JB2CON) were added to each container before starting incubation. The contaminations were added from natural sources (water, glassware, sucrose, pasteurized milk, etc.) to each container. Three containers were prepared for three different temperatures: Ambient room temperature, 35°C, and 44°C. After 21 hours of incubation, microbial contaminants were observed for dahi samples in 100-fold dilution.

For the contamination test, we prepared media and conducted growth promotion tests according to the guideline of the nutrient and dietary supplements section in the United State Pharmacopeia (USP) (Radhakrishna, 2006). Total Non-Lactic Contamination (TNLC) was tested in Tryptic Soy Broth and Tryptic Soy Agar, and Total Yeast and Mold (TYMC) was tested in Tryptic Soy Broth and Sabouraud Chloramphenicol Agar. Moreover, specific pathogens, particularly, Salmonella tested on selective media, such as Rappaport Vassiliadis Salmonella Enrichment Broth, and Xylose Lysine Deoxycholate agar (XLD agar). The presence of *E. coli* was observed in MacConkey Broth and MacConkey Agar. Furthermore, the presence of *S. aureus* and *P. aeruginosa were* checked on Mannitol Salt Agar and Pseudomonas Cetrimide agar respectively (Radhakrishna, 2006).

2.3.1. Analysis of milk for preparation of dahi

Cow milk sample was collected from Dhaka Milk Industry, Dhaka, Bangladesh. Milk was analyzed for four properties, such as lactose, protein, fat, and minerals. Although these four properties were tested in Dhaka Milk Industry and the result was presented in their label, we retested these properties once by a milk analyzer (LACTOSCAN LW, Milkotronic Ltd.) following the method of AOAC (AOAC 13th edition, 2012).

2.3.2. Preparation of probiotic dahi

0.5 L cow milk with 30.0g sucrose was concentrated to 0.4L by heating. Then the temperature of the milk was lowered to about 45°C. 1.0g of the isolate JB2CON was added into each container. The containers were incubated at four different temperature conditions, such as ambient room temperature, 25°C, 35°C, and 44°C for 21 hours. For comparison, the pH of dahi was observed at four temperatures at the set time intervals.

2.4. Shelf-life study at storage condition

For selecting the suitability of the shelf life of probiotic dahi, antimicrobial activity test and shelf-life test were carried out simultaneously. An antimicrobial activity test was performed with some

modifications by the time-kill assay described by Chidre et al. (2017) (Chidre et al., 2017). For this, a total of four containers (each one contained probiotic dahi) was incubated at 44°C for 5.50 hours and then stored at freeze temperature (2°C-8°C) for 14 days. Container-1 was filled with all ingredients of dahi (pasteurized milk, sugar, plus the isolate JB2CON) aseptically, and other containers contained the components of container-1 plus additional contaminations. Container-2 was allowed with unknown contamination (natural) and probiotic isolate, container-3 was filled with *C. albicans* ATCC 10231 plus probiotic isolate, and container-4 was inoculated with *E. Coli* ATCC 8739 plus probiotic

The shelf life of probiotic dahi was tested for 14 days in refrigerated (2°C- 8°C) conditions. *L. acidophilus* must be existed in concentrations of 10⁵ - 10⁶ CFU per mL in order to perform its probiotic actions (Alberts, 1987). As a result, we decided that the dosage of our probiotic dahi for the duration of its shelf life should be at least 10⁶ CFU per mL. The total probiotic count after 14 days was compared with that on the first day at freezing temperature. This comparison was conducted for TNLC, TYMC, *C. albicans* ATCC 10231, and *E. Coli* ATCC 8739. The tests were performed twice.

2.5. Consumer acceptance of dahi

isolate.

Probiotic dahi was prepared according to the process described at point 2.3.2 and incubated at 44°C for 5.50 hours. Consumer acceptance was assessed by a sensory test conducted by the method of Hashim et al. (2009) [24]. 9-point hedonic scale was used, such as, like extremely=9, like very much=8, like moderately=7, like slightly=6, neither like nor dislike=5, dislike slightly=4, dislike moderately=3, dislike very much=2, and dislike extremely=1 (Hashim et al., 2009). 9-point hedonic scale was applied for four groups having four people in each group.

3. Results and Discussion

3.1. Probiotic isolation and identification

3.1.1. Probiotic isolation

In MRS media, several colonies showed lactobacillus characteristics from morphological viewpoints (Figure 1). The colonies were white, round, raised, and translucent (Kavitha et al., 2016).

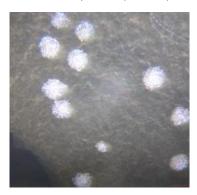


Figure 1. The colony morphology of the colony JB2CON was observed under a stereo microscope, Leica MZ9.5 (Germany).

3.1.2. Gram staining test, 16S gene sequencing, and phylogenetic tree

The cells were gram-positive and long rod (Kavitha et al., 2016). According to the 16S gene sequence, the colony JB2CON was identified as *Lactobacillus acidophilus* (accession no. OM909067, and strain JB2CON). The phylogenetic tree was constructed with the type strains of 10 species in *L. acidophilus* subgroup (Matthew et al., 2013). In the tree of *L. acidophilus* subgroup, our isolated strain JB2CON was placed with AY773947.1 *L. acidophilus* BCRC1069. Therefore, the strain JB2CON was

6

identified as *L. acidophilus*. The tree was rooted with the 16S rRNA gene from AJ276351.1 Bacillus subtilis DSM10 (Figure 2).

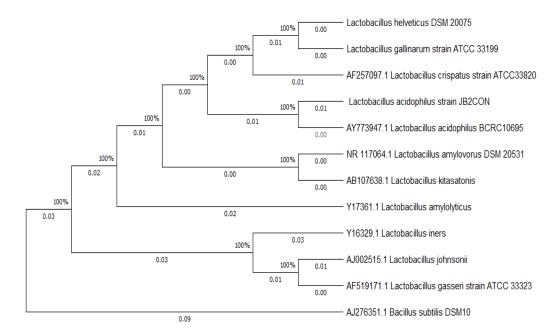


Figure 2. The phylogenetic tree was inferred using the Neighbor-Joining method. The bootstrap consensus tree inferred from 1000 replicates is taken to represent the evolutionary history of the taxa analyzed. Evolutionary analyses were conducted in MEGA11. In the tree with *L. acidophilus* subgroup, our isolated strain JB2CON was placed with AY773947.1. Lactobacillus acidophilus BCRC1069.

2.1.3. Cellular morphology analysis under Transmission Electron Microscope

Under a transmission electron microscope, microorganisms with rod shape and sizes between $2-5 \mu m$ were seen. In the course of our research, we identified rounded-end rods that had been found in pairs, short chains, or even as single cells (Figure 3).

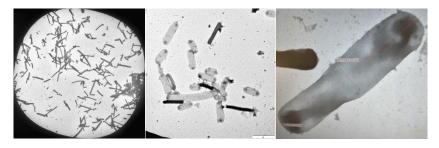


Figure 3. The cellular morphology of *Lactobacillus acidophilus* strain JB2CON in MRS broth was observed under a JEM 1010 electron transmission microscope (JEOL). The left image (1,000 fold magnification) depicted rod shaped free cells and cells in chains. The middle image demonstrated rod shaped cells of varing lentgs and widths at 5000 fold magnification. The right picture showed a single cell of the strain JB2CON at 30,000 fold magnification (length- $4.76 \mu m$ and width- $1.14 \mu m$).

3.2. Probiotic potentiality and safety test

Lactobacillus acidophilus (LA) is the most common probiotic in the world. The probiotic potentiality of these bacteria was studied in previous studies (Kailasapathy and Chin, 2000). In the current study we tested three probiotic features of the isolate, *L. acidophilus* JB2CON (accession no. OM909067).

3.2.1. Antibiotic sensitivity test

L. acidophilus JB2CON (OM909067) in our study had a Zone of Inhibition (ZOI) larger than 20mm (diameter). According to the description of the Clinical and Laboratory Standards Institute (CLSI), they were susceptible to Cefuroxim (100µl/30µg), Gentamycin (100µl/10µg), Chloramphenicol (100µl/30µg), Erythromycin (15 µg / 100 µL), Clindamycin (10 µg / 100 µL), Tylosin (30 µg / 100 µL), Amphicilin (10 µg / 100 µL), Vancomycin (30 µg / 100 µL), Kanamycin (30 µg / 100 µL), Streptomycin (10 µg / 100 µL), and Tetracycine (30 µg / 100 µL) (Table 1) (Prabhurajeshwar et al. 2019, CLSI, 2015). Every test for antibiotic sensitivity was performed thrice.

ZOI Antibiotic Dose $30 \mu g / 100 \mu L$ Cefuroxim 33.0±4.24 $10 \mu g / 100 \mu L$ Gentamycin 22.0±2.0 Chloramphenicol $30 \mu g / 100 \mu L$ 48.67±2.31 $15 \mu g / 100 \mu L$ 31.0±1.0 Erythromycin $10 \mu g / 100 \mu L$ 30.0±6.0 Clindamycin $30 \mu g / 100 \mu L$ 67.33±6.43 **Tylosin** $10 \mu g / 100 \mu L$ 69.33±11.06 Amphicilin Vancomycin $30 \mu g / 100 \mu L$ 48.67±4.16 $30 \mu g / 100 \mu L$ 21.33±2.31 Kanamycin $10 \mu g / 100 \mu L$ 20.67±1.15 Streptomycin $30 \mu g / 100 \mu L$ 42.0±2.0 Tetracycine

Table 1. Antibiotic sensitivity test.

3.2.2. Hemolysis test

Hemolytic activity was not observed for the strain JB2CON. The isolate was identified as γ -hemolytic or non-hemolytic because there was no distinct transparency or greenish zone encircling their colonies on the blood agar Petri-dishes. A probiotic with these qualities is perfect (Halder et. al., 2017).

3.2.3. Acid tolerance

Acid tolerance test was conducted in a solution of pH 2.35, because pH of human stomach in natural state is between 1.5 and 3.5 (Helmenstine , 2020). Probiotic count in the tested solution plunged to $(5.98\pm0) \times 10^3$ CFU/mL after 24 hours from $(1.28\pm0) \times 10^7$ CFU/mL of initial count. Therefore, the probiotic *L. acidophilus* JB2CON (OM909067) was moderately tolerant to pH of gastric juice.

3.2.4. Bile salt tolerance

The good bacteria found in the human colon, known as probiotics, must adjust to bile salt. Since oxgall is typically used for this kind of test, we used it to assess the bile salt endurance of our isolate JB2CON (Hassanzadazar et al., 2012). After incubating the isolate for eight hours, we cultivated it on plates and saw growth on petridishes. When 0.025% and 0.5% oxgall were added, the CFU/mL decreased from (1.37±1.52) ×106 to (1.53±2.14) ×105 and (1.40±2.25) ×104, respectively, while it increased to (2.42±3.11) ×106 in the control broth. The outcomes supported the findings published by Ding et al. (2007), who noted that the probiotic strains in their investigation decreased after eight hours of incubation (Ding et al., 2007). For a bile sensitivity test in MRS broth, Hassanzadazar et al. (2012) selected a number of microbial cultures with bile concentrations as high as 0.3% as the test and 0% as the control. Despite the fact that our study contained a significantly higher concentration of live bacteria than theirs did, this experiment confirmed our findings (Hassanzadazar et al., 2012). As a result, the JB2CON strain is ideal for growing in the bile salt-surrounded human gut.

After 48 hours of incubation, the zone of inhibition was measured. The inhibition zones of more than 20mm, 10-20mm, and less than 10mm were accepted as strong, intermediate, and low inhibition respectively. In our study *L. acidophilus* JB2CON (OM909067) had an intermediate inhibition against tested pathogens (Fijan et al., 2016). We might get a better result than this if we would carry out our experiment with simple spot-on lawn agar, because the Zone of Inhibition (ZOI) of a simple spot-on lawn agar is larger than that of the well diffusion method (Fijan et al., 2016) (Table 2). We found the widest ZOI (14.5 mm) against *E. coli* ATCC 8739, and no ZOI against *C. albicans* ATCC 10231 by agar well diffusion method. That is why we conducted further anti-*E. coli*, and anti-*C. albicans* experiment in time-kill assay method during shelf-life test.

Bacteria Diameter of ZOI (mm)

Salmonella typhimurium ATCC 14028 13.5±0.5

Staphylococcus aureus ATCC6538 12.0±0.0

Escherichia coli ATCC 8739 15.75±0.35

Bacillus subtitlis ATCC 6633 12.25±0.35

Pseudomonas aeruginosa ATCC 1427 12.0±1.41

Candida albicans ATCC 10231 0.0

Table 2. Antagonistic test.

3.2.6. Antimicrobial activity test by time-kill assay for unknown contamination

Total Non-Lactic Contaminations were too numerous to count (TNTC) on TSA media. On the other hand, no CFU of TYMC (Total Yeast and Mold Count) was observed on SCA media at 35°C and 44°C, whereas TNTC was observed at ambient room temperature on this media (Table 3). Further dilution was not performed for counting the exact CFU of unknown contamination, because the tested samples did not comply with the United State Pharmacopeia. In USP nutritional and dietary supplements guideline, microbial limit for TAMC is 1000 CFU/mL, and for TYMC it is 100 CFU/mL (Radhakrishna Tirumalai, 2006a). On the contrary, identical colonies of *E. coli, S. aureus, salmonella spp., and P. aeruginosa* were not found on selective media (Table 4). The samples complied with USP specification in terms of specific pathogenic microorganisms (Radhakrishna Tirumalai, 2006b). It indicates that probiotic dahi prepared by *L. acidophilus* JB2CON (OM909067) can be a natural preservative for preventing the growth of bad bacteria (*E. coli, S. aureus, salmonella spp., and P. aeruginosa*), albeit there are some resistant unknown bacteria and fungus can grow in dahi naturally. Thus, precautions should be taken for preparing dahi and selecting a starter culture.

Types of Types of CFU/mL in CFU/mL in morphology pН Temperature morphology **TSA SCA** in TSA in SCA Ambient room 3 1 4.74 **TNTC TNTC** temperature 2 3.95 35°C TNTC < 100 0 44°C 2 <100 0 3.75 **TNTC**

Table 3. Effect of pH (for certain incubation temperature) on total microbial contaminants.

Table 4. Effect of pH (for certain incubation temperature) on specific pathogenic microbial contaminants.

Microorganism	Media	pH Temperature Identical Colony		
Escherichia coli	MacConkey Broth,	4.74 Am	bient	No Growth
	MacConkey Agar	3.95	35°C	No Growth

		3.75	44°C	No Growth
Staphylococcus aureus	Mannitol Salt Agar	4.74	Ambient	No Growth
		3.95	35°C	No Growth
		3.75	44°C	No Growth
	Rappaport Vassiliadis	4.74	Ambient	No Growth
Salmonella sp.	Salmonella Enrichment Broth,	3.95	35°C	No Growth
	XLD Agar	3.75	44°C	No Growth
Pseudomonas		4.74	Ambient	No Growth
aeruginosa	Pseudomonas Cetrimide agar	3.95	35°C	No Growth
	_	3.75	44°C	No Growth

3.3.1. Analysis of milk for preparation of dahi

Milk quality, such as protein and age of milk influence the titrable acidity of milk, and subsequently, titrable acidity and pH of dahi (Schmidt et al., 1996). Thus, we tested the milk properties of cow milk. Fat content was not less than 2%, and protein content was 3.5%. It was ideal full-fat milk (Table 5) (Maryam et al., 2019).

Serial No. **Ingredients Amount** 1 Fat 2.67 % 2 SNF (Solids-not-Fat) 8.6 % 3 Protein 3.10% 4 Lactose 4.1% 5 Specific gravity 1.027 6 pН 6.81 7 Conductivity 4.1 8 Added Water 2% 9 24.5 Temperature 10 Freezing Point -0.693 11 Salts 0.85%

Table 5. Milk properties.

3.3.2. Preparation of probiotic dahi

The Fastest solidification was found in the case of probiotic dahi at 44°C and it happened within 4.60±0.35 hours when pH reached 4.60±0.35 (Table 6). *L. acidophilus* is very well suited for living in dairy medium, as fermented milk is the ideal method of delivery for introducing *L. acidophilus* into a gut microbiome (Meng et al., 2021).

Table 6. Time, pH, and temperature for formation of dahi.

Temperature	pH of probiotic dahi	Probiotic dahi formation time (hour)	
25	5.55±0	Not completed	
Ambient Room Temperature	4.75±0.01	20.5±0.71	
35	4.85±0.16	4.75±0.35	
44	4.65±0.14	4.60±0.35	

3.4. Shelf-life study at storage condition

A total of four containers (containing probiotic dahi) were incubated at 44°C for 5.50 hours and then stored at refrigerator (2°C-8°C) for 14 days (Table: 16). In the previous study, it was confirmed

that the viability of *L. acidophilus* cells stored at refrigerator (4 °C) is higher than that of cells stored at room temperature (25 °C) (*Arepally et al.*, 2020). Thus, in the current study, we used refrigerator for storing our probiotic.

No *E. Coli* cell was found live on TSA media after 5.50 hours whilst *C. albicans* cells increased gradually and soared to 114 CFU/g after 14 days. This suggests that *L. acidophilus* JB2CON (OM909067) is strongly bactericidal against *E. Coli*. TNLC, and TYMC also increased steadily on TSA and SCA after 14 days. It indicates that probiotic dahi prepared by *L. acidophilus* JB2CON (OM909067) can be a natural preservative for inhibiting the growth of *Escherichia Coli*, although there are some resistant bacteria and fungi which can grow in dahi naturally. Thus, precautions should be taken for preparing dahi and selecting a starter culture. For setting shelf life, these factors should be considered carefully.

The container with pasteurized milk had no microbial contamination. Interestingly, in this container, the probiotic count of dahi increased 2.98-fold and soared to $170.0\pm0 \times 10^9$ CFU/30mL after day 14 from $60\pm4.24\times10^9$ CFU/30mL on day 1. We decided that our probiotic dahi should contain at least 10^6 CFU per milliliter for the duration of its shelf life, in accordance with Alberts' (1987) suggestion (Alberts, 1987). However, our yogurt with probiotics was far more concentrated—up to 14 days.

On the other hand, the pH of dahi went up slightly (0.06) from 3.65±0 to 3.59±0 (Table 9). It indicates that CFU is increasing gradually for 14 days, but pH is not increasing noticeably by this time. This slight change in pH cannot affect the taste (sourness/acidity) of dahi. Therefore, the shelf life of 14 days is the ideal time duration for maintaining CFU and the taste of dahi According to the finding of our study, for maintaining a shelf life of 14 days, probiotic dahi should be prepared from pasteurized cow milk.

Day and pH	Probiotic count (CFU/30mL)	Contamination on TSA (CFU/mL)	Contamination on SCA (CFU/mL)	C. albicans (CFU/mL)	E. coli (CFU/mL)
	Container-1	Container-2	Container-2	Container-3	Container-4
1 & 3.65±0	60±4.24 ×10 ⁹	TNTC	4.5±0.71	94.5±0.71	00
3 & 3.65±0	72.0±0 ×10 ⁹	TNTC	5.0±0	106.0±0	00
14 & 3.59±0	170.0±0 ×109	TNTC	121.0±0	114.0±0	00

Table 9. Time and pH on probiotic bacterial count and contamination count at storage condition.

3.5. Consumer acceptance of dahi

The overall consumer acceptance of dahi was satisfactory (Table 10). 9-point hedonic scale for sensory test of probiotic dahi was applied on total 16 people (four groups). The texture of dahi was neither liked nor disliked by the people of four groups. Basically, texture of dahi depends on various factors, such as, composition of raw milk, incubation time, additives, and so on (Yagmur et al., 2018). In the probiotic dahi, we did not add any extra additive, and fermentation time was reduced to decrease the sourness or acidity of the dahi. Therefore, our prepared probiotic dahi was soft dahi. However, the groups of people liked moderately the overall taste of the probiotic dahi, and also liked slightly or moderately the color, sweet, and sour.

Table 10. Sensory test by 9-point hedonic scale.

Group	Favor	Color	Sweet	Sour	Texture
1	7.25±1.50	6.75±.95	6.0±.81	$6.0 \pm .81$	4.5±.58
2	$7.0 \pm .81$	$7.0 \pm .81$	$7.0 \pm .81$	$7.0 \pm .81$	5.0±0
3	7.5±1.29	7.5±1.29	7.5±1.29	7.5±1.29	5.0±0
4	$7.0 \pm .81$	$6.0 \pm .81$	$6.0 \pm .81$	$6.0 \pm .81$	5.0±0

Note: A 9-point hedonic scale was used where 1 = dislike extremely and 9 = like extremely; mean \pm SD was calculated for 4 people in each group.

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4. Conclusions

The dahi in the extant study was prepared from a potential probiotic, *L. acidophilus* JB2CON (OM909067) isolated from dahi. As the probiotic strain had anti-*Escherichia coli* activity this probiotic, or pribiotic dahi can be an inexpensive option for recovering and preventing diarrheal disease and other *Escherichia coli-related* diseases in humans. This will save globally the cost and lives of billions of people who are infected with *Escherichia coli*. This is the greatest outcome of this research. Furthermore, according to our findings, contaminations in dahi originated from unhygienic practices and unidentified starter culture. These can reduce the shelf life of dahi and cause dahi spoilage. Therefore, precautions should be taken, and dahi can be prepared with pasteurized milk. Considering contamination, probiotic count, pH, and taste of dahi, the time duration of 14 days was an ideal shelf life for our probiotic dahi prepared from pasteurized milk.

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