Amylolytic Capability and Performance of Probiotic Strains in a Controlled Sorghum Fermentation System

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Abstract: This study aimed to explore the fermentative performance of nine lactic acid bacterial strains with probiotic potential during sorghum fermentation. The strain’s attributes including proliferation counts, pH levels, production of organic acid antibacterial activity, and their ability to break down starch were evaluated during the fermentation period in the presence and absence of glucose as a carbon source. In addition, the inhibitory activity of these potential probiotic strains against pathogenic bacteria (Salmonella typhimurium, Escherichia coli, and Staphylococcus aureus) was examined through a co-culturing technique. The results demonstrated that all 4 Lactobacillus strains exhibited robust growth in both glucose and glucose-free fermentation experiments. Glucose supplementation significantly enhanced lactic acid yield which ranged from 0.19 to 0.44% compared to fermentation without glucose which ranged from 0.04 to 0.29%. The selected Lactobacillus strains effectively lowered the media pH below 4.0 after 24 h, producing substantial lactic acid. Notably, in the absence of glucose, only Lb. helveticus D7 and Lb. amylolyticus D12 achieved pH levels below 4 after 8 h, producing the highest lactic acid amounts of 0.27 and 0.29% after 24 h, respectively. Amylase activity was detected on two strains, D7 and D12. Furthermore, most of the tested Lactobacillus strains demonstrated complete inhibition (6 log to 0 Log CFU/mL) of pathogen growth after 24 h of co-culturing, suggesting their potential for enhancing the safety quality of sorghum-based fermented products.

Keywords: fermentation; sorghum; amylase; pathogens; probiotic

1. Introduction

Lactic acid bacteria play a crucial role in many African fermented cereal-based products such as such as Ogi, Mageu, and Togwa by controlling foodborne pathogens. They are used to preserve food, increase the bioavailability of nutrients and functional properties of foods [1–3]. Some LAB species have been well documented for their probiotic properties and have been used in many fermented products such Keffir, Kimchi, and Kombucha, etc., [4]. LAB produces lactic acid as a major metabolic end-product of carbohydrate fermentation and thus exhibits an increased tolerance to acidity. These bacteria contribute to the organoleptic and textual profiles of many fermented foods, and they have been reported to confer health benefits when ingested in adequate amounts [5]. LAB has been extensively used in the fermentation of cereal-based foods in Africa (Kisra, Ogi, Acida, and Merissa). The process of fermentation of carbohydrates by LAB involves the breakdown of six-carbon sugar glucose into two molecules of the three-carbon organic acid, pyruvic acid [6]. This process is coupled with the transfer of chemical energy to the synthesis of adenosine triphosphate (ATP) [7]. Depending on the availability of oxygen, pyruvate can either be oxidized through the tricarboxylic acid cycle or, in the absence of oxygen, produce lactic acid as the major fermentation product [7,8]. Several different substrates (milk, starches, oligosaccharides, glycerol, and other complex carbohydrates) are used for the fermentation of lactic acid by LAB. Therefore, the carbon source for microbial production of lactic acid can either be a sugar in its pure form, or it must be obtained from other substrates, i.e., starch, during fermentation [9].
Amylolytic or amylase-producing LAB (ALAB) plays a vital role in fermentation processes for efficient carbohydrate utilization during the fermentation of starch-based products. ALAB hydrolyzes the starch molecules into polymers composed of glucose units and plays a role in flavor development, preservation, and nutritional enhancement of cereal-based products without the need to add sugar in the fermentation process. Lactic acid fermentation is linked to a decrease in pH with a simultaneous increase in acidity, as with lactic acid and other organic acids. This mechanism results in decreasing the survival time for pathogenic bacteria such as *Escherichia coli* and *Staphylococcus aureus* during food processing [10,11]. LAB have been incorporated into a cereal-based product called togwa [12], and it has been demonstrated to significantly reduce enteropathogenesis in children under 5 years of age.

Sorghum is a staple food for many households and proves to be the grain of importance in Africa due to its high drought tolerance and low price [13]. Sorghum grains contain a substantial amount of carbohydrates, minerals, and vitamins. In Southern Africa, it is consumed in South Africa, Botswana, Zimbabwe, and Lesotho throughout the year [10,13]. LAB isolated from fermented sorghum slurries (Ting) were shown to possess probiotic properties. Similar cereal-based probiotic products include togwa, busheri, etc. Probiotic strains belonging to *Lactiplantibacillus plantarum* are commonly used in the food industry and have frequently been isolated from traditional cereal-based fermentations, and extensive research has highlighted its probiotic potential, showing that *Lb. plantarum* isolated from sourdough can establish itself in the human intestinal tract [14]. The exploration of cereal-based probiotic foods, leveraging the advantages of live cultures emerges as a cost-effective and accessible solution with potential health benefits for diverse populations. Younger children could benefit from high energy, drinkable food products derived from starchy substrate (i.e., sorghum) fermented by amylolytic probiotic strains. These kinds of products could have an added assist in the alleviation of malnutrition and stunting in young children. Malnutrition and stunting are still a problem in most of the developing countries, including South Africa. However, the performance of these probiotics in the preparation of different substrates was not established. In the current study, *Lactobacillus* strains were examined to assess their fermentative capabilities, analyzing microbial counts, amylase activity, pH, and lactic acid production. This study aims to establish the amylolytic activity of probiotic ALAB isolated from Ting (fermented sorghum porridge) and their performance for application in the development of high-energy food products suitable for young children.

2. Materials and Methods
2.1. Bacterial Strains and Culture Conditions

Four probiotic strains were obtained from the Department of Biotechnology and Food Technology's culture collection (Tshwane University of Technology). The strains utilized in the current study had previously been tested for their probiotic potential in the research conducted by [15]. The sequences of these strains were determined and deposited into the National Center for Biotechnology Information database (refer to Supplementary Table S1 for list of strains and their sources). The *Lactobacillus* strains were initially cultivated in a propagation medium to allow them to acclimatize to the fermentation environment. The propagation medium was prepared as follows: MgSO$_4$ (0.5 g/L), yeast extract (5 g/L), dextrose (10 g/L), NaCl (2.5 g/L), KH$_2$PO$_4$ (1.25 g/L), bacterial peptone (1 g/L), calcium carbonate, and sorghum (80 g/L) were carefully weighed and combined in 1 L of deionized water and placed in a cooking pot. The ingredients were heated for 15 min until gelatinization of the sorghum slurry was observed. Subsequently, the slurry was dispensed into 250 mL Schott bottles and autoclaved at 121 °C for 15 min. The propagation medium was inoculated with the probiotic strains listed in Table S1, and the samples were incubated at 37 °C in a water bath for 16 h, to allow the pH of the medium to reach 3.5. This was performed in duplicate, and a volume of 25 mL of the propagated medium was used for each *Lactobacillus* strain for batch fermentation study.
2.2. Amylolytic Activity of LAB Strains

The 4 selected potential probiotic strains were grown on starch agar plates, and starch
was used as the only source of carbon. The assaying method was according to that of [16].
The medium is composed of corn starch (2%), bacterial peptone (0.5%), beef extract (0.5%),
yeast extract (0.2%), sodium chloride (0.2%), and agar (1.5%). A volume of 30 µL overnight
bacterial culture was inoculated at the center of the agar medium. The plates were left to
dry out for 2 h prior to aerobic incubation at 37 °C for 72 h. Thereafter, the colonies were
removed from the agar medium using a sterile blade, and the plates were flooded with
iodine for 30 min. The plates showing a zone of clearing were considered as positive for the
amylase enzyme, and the experiment was repeated twice. Further work was undertaken
to determine the viscosity of fermented sorghum porridge whereby sorghum flour 8% 
(w/v) was mixed with distilled water and cooked for 15 min at 94 °C to make porridge.
The porridge was sterilized using an autoclave, cooled at room temperature and thereafter
inoculated with 4% (v/v) of bacteria strains. The porridges were incubated at 37 °C for 24 h,
and their viscosities were measured using the Brookfield RV viscometer with spindles 4 and
6 at different rotation speeds. All viscosity measurements were performed in thermostated
conditions at 45 °C.

2.3. DNA Amplification of Amylase Gene of Probiotic Strains

Genomic DNA of the probiotic isolates was extracted using a DNA Purification Kit
(United State of America) by following manufacturer’s protocol. DNA amplification
was performed using a thermal cycler (Eppendorf AG 22331 thermal cycler) by target-
ing the amylase gene. The primers used in the study were as follows: amyA -1-F 5′-
CGTCCAGTTGATGGTGGTAATG-3 and amyA -1-R 5′-CCAGGCCAAGCTGAAGTTATAG-
3. The following volumes were measured to perform PCR: 12.5 µL Master mix (KAPA
Biosystems), 1 µL forward primer and 1 µL Reverse primer, 9.5 µL RNase nuclease free
water and 1 µL DNA template to make a reaction volume of 25 µL. The PCR conditions
were as follows: initial denaturation (94 °C for 15 min), 30 cycles of denaturation (94 °C for
30 s), annealing (63 °C for 30 s), extension (72 °C for 3 min), and final extension (72 °C for
7 min).

2.4. Sorghum Fermentation with LAB Strains

2.4.1. Fermentation Medium

The fermentation medium was prepared by combining cooked sorghum (60 g/L)
and glucose (20 g/L), referred to as CSG (cooked sorghum and glucose). The second
fermentation medium solely comprised cooked sorghum (60 g/L) and was designated
as CS.

2.4.2. Population Levels of Probiotic Strains during Fermentation

To determine the population levels of the Lactobacillus strains, 1 mL of each fermen-
tation sample was withdrawn aseptically using a sterile 10 mL syringe. Tenfold serial
dilutions were carried out, and 0.1 mL aliquots were evenly spread on MRS Agar plates
(Merck). The plates were then incubated at 37 °C and bacterial cell counts were recorded at
0, 4, 8, 16, and 24 h.

2.4.3. pH

The pH of the samples fermented with the Lactobacillus strains was measured using
a pH meter (Orion Star-A211, South Africa). The pH meter was calibrated using buffer
solutions at pH 7 and pH 4. Before each analysis, the glass electrode was rinsed with sterile
deionized water. The pH readings were recorded initially and subsequently at 4, 8, 16, and
24 h.
2.4.4. Titratable Acidity

Titratable acidity for each fermented slurry was determined as described by [17]. A volume of 10 mL (diluted 1:10 using distilled water) of each sample was pipetted and titrated against 0.1 N NaOH to phenolphthalein end-point. Three drops of phenolphthalein indicator were added to each diluted sample solution. The concentration of Lactic acid was determined using the equation described below. All measurements were performed in triplicate.

\[
\% \text{ Lactic acid} = \frac{N \times V \times \text{ME of acetic acid}}{\text{Weight of sample in mL}} \times 100
\]

where
- \( N \) is the normality of the sodium hydroxide: 0.1 N;
- \( V \) is the volume of sodium hydroxide (mL) used to reach the titration endpoint;
- \( \text{ME} \) (milli-equivalent of Lactic acid): 0.09008.

2.5. Antimicrobial Activity of LAB Strains Co-Cultured with Pathogenic Strains

Eleven LAB strains were used to determine the antimicrobial activity against indicator pathogenic microorganisms (Escherichia coli ATCC 8739, Staphylococcus aureus ATCC 6538, and Salmonella typhimurium ATCC 14028) during fermentation of sorghum gruels.

The inoculum was prepared by growing LAB strains overnight in 10 mL MRS Broth at 37 °C. Sorghum flour (80 g/L) was mixed with distilled water which was supplemented with glucose. The mixture was cooked for 15 min at 94 °C and sterilized using autoclave; thereafter, the mixture was cooled at room temperature, and the mixture was inoculated with 4% (v/v) of a specific LAB strain together with specific pathogenic (10^6 CFU/mL) strain, and the mixture was allowed to ferment for 24 h in a water bath. Mannitol salt agar was used to enumerate S. aureus, XLD agar for S. typhimurium, and MacConkey agar for E. coli. The microbial counts of pathogenic strains were determined at 0, 4, 8, and 24 h by employing a spread plate technique. The plates were incubated for 24 h at 37 °C. This was performed in triplicate.

2.6. Statistical Analysis

A two-way ANOVA (analysis of variance) test was applied to the experimental data, and the means were compared by Fisher’s test at \( p < 0.05 \) using the software Statgraphics Centurion XVI version 16.1.11 (Statpoint, Warrenton, VA, USA).

3. Results

3.1. Amylase Assay of Probiotic Strains

The amylase activity of the nine LAB strains isolated from household fermented sorghum slurries were grown on starch agar plates, and starch was used as the sole source of carbon; the plates showing zones of clearing were considered positive for amylase enzyme. Supplementary Table S2 shows the results for the amylase assay of the selected LAB strains. Only two strains, D7 and D12, tested positive for the amylase enzyme as compared to other strains.

Figure 1 shows the effect of amylolytic lactic bacteria on the viscosity of sorghum porridges. The two tested Lactobacillus strains D7 and D12 were able to decrease the viscosity of the sorghum porridges during fermentation over 24 h of incubation as shown in Figure S1 (refer to Supplementary Material). Strain D7 decreased the viscosity of fermented sorghum porridges from 2854 to 1209 cP after 24 h, and strain D12 decreased the viscosity from 2812 to 940 cP as compared paired to the controls (K20 and T12). Strain D12 was the best strain in reducing the viscosity of the sorghum porridge with 940 cP after 24 h of incubation as compared to strain D7. This implies that the effect of amylolytic Lactobacillus strains on viscosity could be ascribed to direct effects of the amylase enzyme from amylolytic activity of LAB from the fermentative liquification of sorghum porridges.
The amylase gene was further confirmed through PCR amplification for strains D7 and D17 which were successfully amplified. The results of PCR amplicons for the detected gene are shown in Supplementary Figure S2.

3.2. Population Levels of Lactobacillus Strains during Fermentation in the Absence and Presence of Glucose

The results of two individual fermentation media, one containing glucose and the other without, are presented in Supplementary Material (Tables S3 and S4). In the presence of sugar, all Lactobacillus strains exhibited strong growth, sustaining colony counts between 7 and 8 log10 cfu/mL following an 8 h incubation period. Specifically, strains K20 and D7, showed colony counts within the range of 8.2 to 8.01 log10 cfu/mL. On the other hand, strains D12, and T12 exhibited mean colony counts within the range of 7.5 to 7.6 log10 cfu/mL. Significantly higher colony counts were observed at the end of the 24 h fermentation period. Notably, the following strains exhibited the highest colony counts, ranging from 8.2 to 8.8 log10 cfu/mL: K20, T12, and D7.

Interestingly, the absence of glucose did not significantly impact the growth of the tested Lactobacillus strains (refer to supplementary Table S2). At the 8 h incubation period, strains K20 and T12 displayed the highest mean colony counts, ranging from 8.5 to 8.6 log10 cfu/mL. They were followed by strains D7 and D12 which had mean colony counts ranging from 7.2 to 7.4 log10 cfu/mL. After 24 h of incubation, all the strains had the exhibited highest colony count ranging from 8.2 to 8.6 log10 cfu/mL. Remarkably, as illustrated in Figure S1, virtual liquefaction observations were made on Lactobacillus strains D7 and D12 at the end of the fermentation, as they successfully liquefied the fermentation medium without the addition of glucose.

3.3. pH

Figure 2 illustrates the pH of the fermented sorghum medium in the absence of 10% glucose by selected LAB strains. It was evident that some strains failed to decrease the pH of the fermentation medium to the required level of pH 3 in the absence of glucose. At the 4 h interval, there was already a slight decrease in pH for the tested strains. D12 had managed to decrease its pH to 3.93 as compared to other strains (D7, K20, and T12) which had a pH ranging from 4.0–4.4. At 8 h of incubation, strain D12 further reduced its pH to 3.7, which was followed by D7, which decreased its pH from 4.1 to 3.9, and the remaining strains had minimal pH reduction. The two Lactobacillus strains D7 and D12 continued to

Figure 1. Viscosity analysis of fermented sorghum porridges with strains K20, T12, D7 and D12 over 24 h. Bars with the same letter are not significantly different ($p > 0.05$). Values reported are means of at least three independent measures.
further reduce in pH to 3.5 for both strains at the end of fermentation, followed by T12, with a pH of 3.9. Strain K20 failed to reduce the pH of the fermented sorghum to below pH 4.

![Bar graph showing pH levels of sorghum gruel fermented with LAB strains.](image)

**Figure 2.** The pH for sorghum gruel fermented with LAB strains in the absence of 10% glucose. pH levels were monitored hourly throughout the fermentation process. Values reported are means of at least three independent measures. Bars with the same letter are not significantly different (p > 0.05).

Figure 3 also illustrates the pH levels of *Lactobacillus* strains at various time intervals in the presence of added glucose. The addition of glucose to the fermentation medium provided a competitive advantage to the selected *Lactobacillus* strains. This advantage was particularly noticeable at the 4 h mark, as all the strains (D12, D7, K20, and T12) had successfully reduced the pH below 4; the pH of the samples were in a range of 3.88 to 3.74. At the 8 h period, all the probiotic strains continued to further decrease the pH. By the end of the 24 h fermentation period, all the strains had achieved a pH lower than 4.

![Bar graph showing pH levels of sorghum gruels fermented with LAB strains in the presence of 10% glucose.](image)

**Figure 3.** The pH for sorghum gruels fermented with LAB strains in the presence of 10% glucose. pH levels were monitored hourly throughout the fermentation process. Values reported are means of at least three independent measures. Bars with the same letter are not significantly different (p > 0.05).
3.4. Titratable Acidity

The lactic acid concentration (% w/v) for both fermentation batches, one with the addition of glucose and one without, is illustrated in Table S5 (Supplementary). In this investigation, the fermentation batches that were inoculated with the Lactobacillus strains (K20, T12, D12, and D7) in the absence of glucose exhibited lactic acid concentrations ranging from 0.18% to 0.48% after a 24 h period. Notably, the strains that demonstrated the highest production of lactic acid were D12 and D7, with lactic acid concentrations of 0.48% and 0.42%, respectively, followed by strains K20 and T12 with lactic acid concentrations of 0.18% and 0.24%, respectively.

In contrast, when glucose was introduced, all the Lactobacillus strains demonstrated a competitive advantage, resulting in significantly higher lactic acid concentrations in comparison to fermentations without glucose. The strain that produced the highest lactic acid was T12 and D7, which had a percentage range of 0.67% and 0.68%, respectively, and this was followed by strain K20 and D12 which had a lactic acid percentage of 0.44% and 0.55%, respectively. Strains with the lowest lactic acid production included K20, J20, J22, and V, with concentrations ranging from 0.41% to 0.44%. The results suggest that the addition of glucose significantly increased lactic acid production compared to the batch without glucose. However, in the absence of glucose, strains (T12 and K20) faced challenges in achieving high lactic acid levels, with the notable exceptions of strains D7 and D12 which might contain the amylase gene.

3.5. Antibacterial Activity of LAB Strains against Pathogenic Strains in Sorghum Gruel

The inhibition results of the three pathogenic strains during the co-culturing assay with LAB are represented in Figures 4–6. E. coli was inhibited by all the tested LAB strains; it was noted that the growth of the E. coli strain co-cultured with test LAB strains in sorghum flour was below that of the control experiment from 12 to 24 h of incubation (Figure 4), meaning that there was a suppression of the E. coli strain. Total inhibition was observed with strains (K20, D12, and D7) after 24 h of incubation as compared to strain T12 which inhibited E. coli growth from 7.6 Log CFU/mL to 3.14 Log CFU/mL.

Figure 4. Antibacterial activity of the 4 tested LAB strains against E. coli ATCC 8739 in sorghum gruel fermentation. Bars with the same letter are not significantly (p > 0.05) different. Values reported are means of at least three independent measures.
compared to strain T12 which reduced *S. aureus* growth from 6.6 to 2.3 Log CFU/mL. Values reported are means of at least three independent measures.

All the tested LAB strains were able to inhibit *S. aureus* (T12, D12, and D7). However, after 24 h, total inhibition was observed from the strains as compared to strain T12 which reduced *S. aureus* growth from 6.6 to 2.3 Log CFU/mL. Values reported are means of at least three independent measures.

Figure 5 shows the inhibition pattern of *S. aureus* growth against individual LAB strains. After 12 h of incubation there was slight inhibition of *S. aureus* by LAB strains (T12, D12, and D7). However, after 24 h, total inhibition was observed from the strains as compared to strain T12 which reduced *S. aureus* growth from 6.6 to 2.3 Log CFU/mL.

Figure 6 below illustrates the antibacterial activity of LAB strains against *Salmonella typhimurium* ATCC 14028. All the tested LAB strains were able to inhibit *S. typhimurium* after 24 h of incubation. Total inhibition of the pathogen was observed with LAB strains K20, T12, D12, and D7.
4. Discussion

Lactic acid fermentation remains the preferred method for food processing due to its ability to preserve foods, enhance shelf life, and improve flavor [18]. Cereals fermented with LAB have been proven as a good vehicle for delivering beneficial bacteria. The applicability of probiotic strains in functional food development requires to survive processing conditions and low pH derived from lactic acid production [17–19]. In this study, sorghum was chosen as a vehicle for the selected probiotic strains and as a climate smart crop and is also a cheaper commodity for many Southern African households. The four selected probiotics were subjected to controlled fermentation to assess amylase activity, microbial counts, pH, and lactic acid levels in the absence and presence of glucose. Furthermore, the selected strains were also tested for inhibitory activity against pathogenic bacteria.

There were differences between the two fermentation batches with regards to pH values which made it clear that, in the presence of sugar, the tested *Lactobacillus* strains were all able to acidify the fermentation medium to pH levels below 4. This was attributed to the readily available sugar (glucose) which can be easily used by *Lactobacillus* strains in the fermentation medium because of the observed accelerated growth. In the absence of glucose, it was seen that most of the *Lactobacillus* strains tested were struggling to reduce the pH to below 4. Only three strains, D12, D7, and T12 were able to reduce their pH to below 3. Strains D12 and D7 had the lowest pH of 3.5 as compared to T12 which had a pH of 3.9. It could be possible that the amylase gene detected in our earlier study from the two strains, D12 and D7, had a direct role during fermentation by breaking down starch into simple sugars which can be easily utilized by LAB [20]. Hence, the two strains D12 and D7 were able to reduce the pH to below 4 after 8 h of incubation as compared to other strains. Detection of amylase activity in LAB has been demonstrated by several researchers [20,21]. The amylase enzyme in LAB has been reported to play a direct role on starch molecules by converting starch into simpler monomers which aid liquefaction of starch-based products [22]. Addition of amylase-producing LAB in fermentation of cereal-based products will aid in increasing the fluidity of porridges and increase energy density of cereals such as maize, sorghum, and millets for infants. This study is in line with the work conducted by [22–24] on the amylolytic effect of LAB in cereal-based fermentation.

Furthermore, a decrease in pH is a result of *Lactobacillus* strains being able to convert a simple sugar (glucose) into lactic acid ($\text{C}_6\text{H}_{12}\text{O}_6 \rightarrow 2\text{C}_3\text{H}_6\text{O}_3$), which resulted in a low pH of the fermentation medium. The result of the present study is in agreement with the work conducted by [22,25], who reported a significant pH decrease in the fermentation of sorghum and maize sorghum flours [16,26] also reported production lactic acid and other organic acids which included acetic acid, formic acid, and succinic acid in the fermentation of sorghum flour by *Lactobacillus* species. A reduction in pH and increase in LAB followed a trend reported by [27]. Moreover, a study conducted by [17] has demonstrated that the addition of glucose to the fermentation medium has a direct link to higher organic acid production, which is in line with our current study. The Lactic acid values recorded in this study were in line with other studies [27–29].

Taking into consideration the addition of sugar in product quality and development plays a significant role due on the growth and metabolic activity of the *Lactobacillus* strains, and it may yield products with better quality attributes such as improved taste, texture, and shelf life. However, in regions where sugar is expensive and difficult to procure, it becomes crucial to explore alternatives such as using amylolytic *Lactobacillus* strains that can break complex starches into simpler sugar which can be used by the *Lactobacillus* strain to produce similar attributes as observed in fermentation conducted with the addition of sugar. In the current study, *Lactobacillus* strains D12 and D7 performed better in fermentation without added sugar and is in line with a study conducted by [20] who reported higher lactic acid production using amylolytic *Limosilactobacillus fermentum* 04BBA19 [30]. Furthermore amylolytic *Lactobacillus* have been reported by other studies from different tropical starchy fermented foods such as cassava and sweet potato or grains as maize [31].
Antibacterial activity of 4 probiotic strains (K20, D7, D12, and 12) were tested on clinically important food pathogens, namely, *E. coli* ATCC 8739, *S. aureus* ATCC 6538, and *S. typhimurium* ATCC 14028 by co-culturing with the selected probiotic strains in the fermentation medium. The LAB strains that were able to completely inhibit all the three indicator pathogens were D12 and D7. Strain K20 only inhibited *S. typhimurium* and *E. coli* completely, and strain T12 only inhibited *S. typhimurium* and *S. aureus*. The inhibition of the three pathogens is attributed to the production of fermentation byproducts such as lactic acid, acetic acid, and small portions of propionic acid. These metabolites synergistically work to inhibit the growth of many target pathogens by making the environment unfavorable for most pathogens by reducing the pH to acidic levels [26,31,32].

*Escherichia coli* is classified as gram-negative bacteria within the Enterobacteriaceae family that colonizes the human gastrointestinal tract but has the potential to induce intestinal or extraintestinal infections [33,34]. These infections cause severe invasive diseases, including bacteremia and sepsis. Notably, *E. coli* stands out as the most common cause of bacteremia in both low- and high-income settings, surpassing other pathogens like *Staphylococcus aureus* and *Streptococcus pneumoniae*. Additionally, *E. coli* is a principal causative agent of neonatal meningitis [15,34]. Therefore, consumption of a fermented sorghum beverage may also aid in therapeutic management of various clinical infections and foodborne diseases attributed to *E. coli*, *S. typhimurium* and *S. aureus*.

It can be deduced from the observations made in Figures 4–6 that the use of one strain in the preparation of fermented porridge/products will not be effective in inhibiting enteropathogens [35]. The use of combing probiotic strains for the preparation of fermented beverages would help to prevent disease causing foodborne pathogens such as *E. coli*, *S. aureus* and *S. typhimurium*. Therefore, combing strains that showed total inhibition with the strains that showed little to moderate inhibition will ensure the safety of fermented sorghum beverage. The findings of the current study are in line with the work conducted by [35,36]. The study conducted by [36] reported antibacterial activity of *Lb. plantarum* which completely inhibited the *E. coli* D1288 within 8 h of incubation during the fermentation of Ghanaian maize dough. Several studies conducted on fermented cereal-based porridge (maize dough, motoho, and ting) have demonstrated antimicrobial action against a variety of diarrheal pathogens [10,36].

5. Conclusions

This study focused on assessing the performance of probiotic strains in a sorghum-based medium. During fermentation, all selected *Lactobacillus* strains demonstrated robust growth, maintaining levels above 6 log10 cfu/mL after 24 h, regardless of the presence or absence of sugar. However, fermenting without the addition of glucose resulted in a limited reduction in pH, ranging from 4.0 to 4.4 at the end of the 24 h incubation. Notably, strains D7 and D12 with amylolytic activity achieved a faster reduction, reaching pH 3 within 8 to 24 h. Regarding lactic acid production, strains D7 and D12 exhibited higher levels, measuring 0.42 and 0.48 units, respectively, as determined by titratable acidity. With regards to antibacterial activity, strains D12 and D7 successfully inhibited *E. coli*, *S. aureus* and *S. typhimurium* to undetectable levels after 24 h, while strains T12 and K20 did not inhibit all the three indicator pathogens to undetectable levels after the same duration. These findings suggest the potential for employing combined probiotic strains in the future development of sorghum-based beverages for impoverished communities who cannot afford probiotic supplements.

**Supplementary Materials:** The following supporting information can be downloaded at: https://www.mdpi.com/article/10.3390/fermentation10060308/s1, Figure S1: Fermentation products showing the effect of amylase positive strains D7 and D12 on sorghum gruel; Figure S2: Agarose gel electrophoresis showing positive results for PCR amplification of amylase gene in strains D7 and D12; Table S1: Molecular identification of LAB isolates based on the 16S rRNA and *phe* gene sequencing; Table S2: Amylase assay of the 4 probiotic strains; Table S3: Colony counts of LAB strains used for fermentation over period of 24 hours supplemented with 10% glucose; Table S4:
Mean log colony counts of LAB strains used for fermentation over period of 24 hours without added glucose; Table S5: Lactic acid composition data for fermented sorghum gruels. Percentage lactic acid levels were monitored hourly throughout the fermentation processes. The results were recorded as mean values.

**Author Contributions:** S.M.R. and M.-L.T.-Z. conceived and designed the experiments; S.M.R. performed the experiments; S.M.R. and M.-L.T.-Z. analyzed the data; S.M.R. prepared the first manuscript draft. M.-L.T.-Z. reviewed and edited the final manuscript. All authors have read and agreed to the published version of the manuscript.

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**Data Availability Statement:** All relevant data generated in this study are available in this manuscript and the Supplementary Materials.

**Conflicts of Interest:** The authors declare that there are no conflicts of interest.

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