

Original Research Article

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## Probiotic Characterization of Lactic Acid Bacteria Isolated From Local Fermented Vegetables (Makdoos)

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### ABSTRACT

The present study aims at isolation and characterization of *Lactobacillus* species from Makdoos a locally stuffed and fermented vegetables and to characterize some selected isolates for their probiotic potential. Sixteen isolated *Lactobacillus* strains (7 *Lactobacillus plantarum* 1, 3 *Lactobacillus pentosus*, 1 *Lactobacillus brevis* 1, 1 *Lactobacillus brevis* 3, 1 *Lactobacillus rhamnosus*, 1 *Lactobacillus salivarius* and two were not fully assigned to a species were chosen for further testing. These were tested for their probiotic potential. This included survival at low pH and in gastrointestinal simulated juice, antagonistic activity against *Salmonella typhimurium*, *Escherichia coli*, *Bacillus cereus* and a methicillin-resistant *Staphylococcus aureus* (MRSA) isolate, bile tolerance, antibiotic resistance to 8 antibiotics and adhesion to Caco-2 cells. Most isolates, especially *Lactobacillus plantarum* 1, were tolerant to acidity and intestinal conditions after exposure for three and four hours respectively with reduction less than one log cycle of the starting CFU/ml. The same trend was observed in respect to bile tolerance with slight variations. All isolates inhibited the growth of the tested pathogens and were highly effective against *Bacillus cereus*, *Salmonella typhimurium* and MRSA isolate. As for antibiotic resistance, it was pronounced against tetracycline, streptomycin, kanamycin, and trimethoprim. Some isolates, M5 and M6 showed resistance to 6 or more of the 8 antibiotics tested and others were resistant to 3 or 4 antibiotics. As for adhesion ability to Caco-2 cells, some isolates showed superior ability i.e more than 80% relative adhesion compared with *Escherichia coli* and higher than the probiotic strain *Lactobacillus reuteri* DSMZ 20056. In Conclusion, the ability of these isolates to survive well under the tested conditions as well as being non-hemolytic renders them as good probiotic candidates for health and food technology benefits. Results also prove that the traditionally fermented vegetables are good sources for probiotic *Lactobacillus*. However, further *in vivo* studies are needed to substantiate their potential use in different applications.

#### Keywords

Lactobacillus,  
Makdoos,  
Fermented  
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### Introduction

In the last few years, probiotics use for different puposes had aroused interest in different medicinal settings in different countries (Angmo *et al.*, 2016). Lactic acid

bacteria especially *Lactobacillus* species have been explored as probiotics due to the growing evidence of their health benefits (Feng *et al.*, 2015; Tropcheva *et al.*, 2014).

Sources for the isolation of unique probiotic isolates vary according to geographic regions and dieting habits, however, human and animal gastrointestinal tracts played a great role in isolating promising probiotics (Lieven-Le *et al.*, 2014). In the last decade new resources have been tested including dairy products (Carafa *et al.*, 2015), fermented meat products (Rubio *et al.*, 2014), foods of plant origin (Wang *et al.*, 2014) and traditionally fermented and pickled vegetables (Mahasneh *et al.*, 2015). Earlier studies were mainly concerned in the search for new starter cultures for different fermentations in food production and technology. Entering the probiotics era necessitates characterization of these bacteria for the putative health benefits they provide, as well as searching traditionally fermented vegetables for superior probiotics. Characterization of such bacteria isolated from fermented vegetables is rather sparse and it focuses mainly on their antimicrobial activity. Few studies investigated probiotic attributes of bacteria of pickled and fermented vegetables. As a result more information is needed to substantiate the role of fermented vegetables derived probiotics through studying their adhesion ability, cholesterol reduction, resistance to antibiotics and resistance to acid and bile and other characteristics both in vitro and in vivo.

The increased demand for complimentary and health food encourages innovation and new and novel product development in the food industries (Guo *et al.*, 2010; Abbas *et al.*, 2014). It is well established that consumption of probiotic bacteria as formulation or fermented food ingredients stimulates growth of beneficial bacteria and reduces pathogen activity (Chiang *et al.*, 2012). To attain the health benefits of probiotic foods, it should contain no less than  $10^7$  CFU of viable bacteria per gram. In this context, probiotic fermented or fortified foods have received wide interest in an expanding market (Argyri

*et al.*, 2013). For *Lactobacillus* strains to exert the expected benefits as probiotics, they should fulfill all or most of the criteria laid down for probiotics in general including ability to survive in the gastrointestinal tract, tolerate bile salts as well as adhesion and persistence to resist pathogens through production of antimicrobial substances and other means (Lee *et al.*, 2011; Tulini *et al.*, 2013).

Lactic acid bacteria mainly *Lactobacillus* and *Bifidobacterium* species are considered very basic in probiotic development, however lactobacilli are the fundamental group. They thrive in naturally fermented or fortified dairy products. Recently, a drive towards non-dairy novel probiotics is observed to span traditionally fermented foods of vegetable origin, (Feng *et al.*, 2015). No doubt such traditionally fermented foods would be a unique mining area for new and novel probiotic *Lactobacillus* isolates. It is recognized that wild probiotic strains would compete better in food traditional fermentation settings. Accordingly, new and novel specific probiotic candidate bacterial strains are being sought. The efficacy of such strains is mandatory and should be carefully evaluated for safety.

In Jordan, pickled and traditionally fermented vegetables form a reasonable part of the home-made stored foods. Makdoos in its different forms, is an example, it is made traditionally of fermented aubergine stuffed with ground walnuts, garlic, hot dried pepper and left to ferment in vegetable oil, mainly olive oil (24.8) The other type of Makdoos is the big green pepper Makdoos which is stuffed with cut tomato, parsley, garlic and hot dried pepper and traditionally soaked and fermented in vegetable oil preferably olive oil or a mixture of oil and water. The fermentation process starts with a wide variety of indigenous microorganisms present

in / on the vegetables and the stuffing material used. However, the bacteria responsible for the major fermentation in this case are lactic acid bacteria mainly *Lactobacillus* spp.

Due to increasing concern of the interrelationship between diet and health, great attention is given to the functional properties of indigenous lactobacilli involved in traditional fermented foods (25.13.5.26). It is assumed that these foods could provide a rich source of new novel probiotics with unique properties. The aim of this study was to isolate and assess the pro biotic potential of some *Lactobacillus* species from Makdoos through studying some characteristics such as tolerance to gastrointestinal juice acid and bile), their antagonistic activity against some pathogens, their antibiotic resistance as well as studying their adhesion to Caco cells.

## **Materials and Methods**

Collection of fermented vegetable samples (Makdoos) and bacterial growth enrichment.

Homemade and retail commercial samples were collected from Amman markets during the period 2013-2014 and included in this investigation. These samples of Makdoos were either single samples of stuffed and fermented vegetables or composite samples prepared when needed irrespect of their number. The enrichment process was carried out by inoculating approximately 1 ml of a sample homogenate into 50 ml sterile MRS broth (Oxoid, UK) and incubated anaerobically at 37°C for 2-5 days (13). All samples were collected into sterile glass samplers and were kept in the laboratory at room temperature for further analysis.

### **Isolation of *Lactobacillus* strains**

Enriched Makdoos samples were serially diluted in sterile normal saline. Aliquots of

100 µl from each dilution were then plated onto de Man, Rogosa and Sharpe agar (MRS, Oxoid, UK) supplemented with 0.01% bromocresol purple as a pH indicator. Plates were incubated anaerobically using anaerogen bags (AnaeroGen, UK) at 37°C for 48 hours. Presumptive *Lactobacillus* colonies with yellow halos were randomly picked off the MRS plates and were further subcultured onto fresh plates of the same medium to ensure purity.

### **Identification of bacterial strains**

All isolates were tested for catalase and oxidase activity, Gram reaction, cell morphology as well as spore formation (27.28). The strains were tested for production of acids from carbohydrates and related compounds using API 50 CH kits and CHL media (BioMèrieux, France) according to the manufacturer's instructions. Results were scored after incubation at 37°C for 24 and 48 hours. These results were joined to the apiweb™ identification software with database (V5.1) which uses the phenotypic data to predict a species identity.

Interpretations of the fermentations profiles were facilitated by comparing all results obtained for the tested isolates with information from the computer aided database. Isolates were also tested for their hemolytic patterns, gelatinase and DNase activity according to (Gupta *et al.*, 2007).

### **Maintenance of bacteria**

Bacterial cultures were maintained in MRS broth with 20% glycerol and kept stored at -80°C. Working cultures were kept on MRS agar plates at 4°C and were routinely subcultured every 2-4 weeks. For comparative purposes, *Lactobacillus reuteri* DSMZ 20056, a probiotic strain, was included in some tests.

### **Preparation of simulated gastric and intestinal juices**

Fresh simulated gastric and intestinal juices were prepared daily by suspending pepsin (P7000-25G Sigma-Aldrich, USA) at 0.3% w/v and pancreatin USP (P-1500, Sigma-Aldrich, USA) at 0.1% w/v in sterile 0.5% w/v NaCl. The pH was adjusted to 3.0 for gastric juices using HCl and pH 8.0 for intestinal juices with 0.1M NaOH using pH meter (Eutech 510, Singapore).

### **Bacterial tolerance to simulated gastric and intestinal juices**

Overnight bacterial cultures grown in MRS broth of each test isolate were adjusted to 0.5 McFarland standard (to know the initial bacterial counts per ml) and 30 ml aliquot of that suspension was centrifuged (2500 x g, for 20 minutes, at 5°C), washed twice in 50 mM K<sub>2</sub>HPO<sub>4</sub> (pH 6.5) and resuspended in 3 ml of the same buffer. One milliliter of each isolate suspension was harvested by centrifugation (12,000 x g, for 20 minutes, at 5°C) and resuspended in 9 ml of gastric solution.

Total viable counts on MRS plates were recorded, both before and after incubation period of 3 hours at 37°C. Then, one milliliter of gastric juice was taken and added to 9 ml of intestinal juice solution.

Total viable counts on MRS plates were also recorded, after an incubation period of 4 hours at 37°C. The results were expressed as colony forming units (log<sub>10</sub> orders CFU/ml).

### **Determination of total viable counts**

Total viable counts of *Lactobacillus* species were determined by spread plate method using MRS agar. Serial ten-fold dilutions were prepared using sterile normal saline. Triplicate plates of each suitable dilution were

inoculated with 100 µl each and incubated anaerobically (AnaeroGen, UK) at 37°C for 48 hours after which numbers of CFU/ml were determined.

### **Detection of antibacterial activity of the bacterial isolates**

For the detection of antagonistic activities of the isolates, an agar spot procedure was used. The antibacterial activity of the selected *Lactobacillus* isolates was determined by the test described by (13.30) with some modifications as follows: five microliters of each overnight culture of *Lactobacillus* isolate grown in MRS broth were spotted onto the surface of MRS agar plates (containing 0.2% glucose) and were then incubated under anaerobic conditions at 37°C for 48 hours. An overnight culture of four indicator strains (*E. coli* ATCC 25922), (*S. typhimurium* ATCC 14028), (*B. cereus* toxigenic strain TS) and (MRSA clinical isolate) were grown in nutrient broth and were adjusted to 0.5 McFarland solution standard which is equivalent to about 10<sup>8</sup> CFU/ml. Aliquots of 0.25 ml were inoculated into 7 ml of soft/semi-solid nutrient agar (containing 0.2% glucose and 0.7% agar). Inoculated semi-solid agar was immediately poured in duplicates over the MRS plate on which the tested *Lactobacillus* isolate was grown. The plates were incubated aerobically to allow growth of the test pathogenic bacteria at 37°C for 24 hours. The antibacterial activity was detected by measuring the inhibition zones around the *Lactobacillus* bacterial spots. Inhibition was recorded as positive if the diameter of the zone around the colonies of the producer was 2 mm or more (Abbas *et al.*, 2015). This result were further confirmed by testing the neutralized cell free culture supernatant by the agar diffusion assay to eliminate the probability of inhibition by the acid produced by isolates.

### **Bile tolerance test**

The tolerance of the bacterial isolates to bile was tested using MRS broth prepared with 0.3, 0.5, 1 and 2% (w/v) oxgall (Oxoid, UK). Ten milliliter aliquots of bile solutions were transferred into standard test tubes and sterilized by autoclaving at 121°C for 15 min. Bacterial cultures were inoculated into sterile MRS broth, incubated overnight and adjusted to 0.5 McFarland turbidity standard at the time of use. Two hundred microlitres of the adjusted bacterial cultures were inoculated into different bile concentrations tubes for each isolate. One milliliter aliquots were taken from each inoculated bile tube at zero hours of incubation and after 24 hours, serially diluted with sterile normal saline and spread plated in triplicates onto MRS agar plates to determine total viable counts.

### **Antibiotic susceptibility testing**

The antibiotic susceptibility test was done according to the agar diffusion method published by the National Committee for Clinical Laboratory Standards (NCCLS, 2000). The determination of minimum inhibitory concentration (MIC) to certain antimicrobial agents recommended by Scientific Committee on Animal Nutrition included Ampicillin, Ciprofloxacin, Erythromycin, Gentamicin, Kanamycin, Streptomycin, Tetracycline and Trimethoprim. Müller-Hinton agar (Merck, Darmstadt, Germany) plates were used and incubated under anaerobic conditions. Serial dilutions of antibiotics were prepared using sterile distilled water, DMSO and/or ethanol and were sterilized using 0.22 µm syringe filters (Macherey-Nagel, Germany). One ml of each suitable antibiotic concentration was added to 9 ml of molten agar, mixed thoroughly and poured into sterile petri dishes. The agar plates were allowed to set at room temperature. Bacterial inocula were

prepared by suspending several bacterial colonies from fresh agar plates in normal saline to a 0.5 McFarland turbidity standard. A spot of 4 µl of the inocula was placed on the agar surface. The inoculated plates were allowed to stand at room temperature for about 30 minutes. The triplicate plates were then transferred into anaerobic jars and were then incubated at 37°C for 24 hours. The MIC (Minimum Inhibitory Concentration) was recorded as the lowest concentration of the antimicrobial agent that completely inhibited growth.

### **Adhesion to Caco -2 cells**

The test was conducted based on the method described by Balamurugan *et al.*, (2014). To grow the Caco-2 cells, 10 percent v/v inactivated fetal calf serum, 1 percent non-essential amino acids, 1 percent glutamine and 20 microgram per ml of streptomycin and penicillin were added to Dulbecco's modified Eagle medium (DMEM). Caco-2 cells were added to this medium. Seeded into 24-well tissue culture plates at a concentration of 1 million cells per ml and incubated at 37 C in 10 percent CO<sub>2</sub> for seven days.

Medium was changed every two days. Bacterial isolates were grown in MRS broth, pelleted and washed and finally suspended in non-supplemented DMEM to a concentration equal to 3 McFarland standard. One ml of this suspension was overlaid onto caco-2 cells monolayers and incubated at 37C in 10 percent CO<sub>2</sub> for 90 minutes. The bacterial suspension was then removed via aspiration, DMEM was used to wash the cells and one ml of 0.1 % Triton X-100 was added for 10 minutes to detach any bacterial cells that have adhered to the Caco -2 monolayer. The detached bacterial cells were then plated onto MRS agar at 1:100 and 1:1000 dilutions and then incubated at 37 C in 10% CO<sub>2</sub> for 24 hours. Colonies were then counted after



incubation and *E. coli* was used as a positive control i.e 100 percent relative adhesion.

### Statistical analysis

The results are presented as means  $\pm$  SD. Statistical differences among bacterial isolates were determined by two way ANOVA except for tolerance to simulated gastric and simulated intestinal juices which were determined by three way ANOVA. Differences were considered significant at  $p < 0.05$ .

## Results and Discussion

### Isolation and identification of *Lactobacillus* potential probiotic strains

A total of sixteen isolates from different fermented samples were chosen to be used in this study. All isolates were Gram positive rods, catalase and oxidase negative, non-spore forming, non-hemolytic, and DNase as well as gelatinase negative (Table 1). Absence of hemolytic activity of these isolates is a positive sign in favour of being suitable probiotic isolates irrespective of being *L. plantarum*, *L. pentosus*, *L. brevis* or *L. salivarius*. Similar observations were recorded for *Lactobacillus* isolates from dairy and other sources. These isolates were further characterized using API 50 CH strips. Results of the API 50 test confirmed the identity of the 16 *Lactobacillus* isolates (Table 2). Identification of the isolates (Table 2) indicated the dominant presence of *Lactobacillus plantarum* where 7 out of the 16 isolates belonged to this species, followed by 3 *Lactobacillus pentosus* isolates and 2 *Lactobacillus brevis*. One each of *L.rhamnosus*, *L.salivarius*, and two isolates were not designated to any species.

Probiotic lactobacilli have been isolated from foods of plants origin and cereals. It is recognized that wild type strains that

dominate naturally fermented products tend to have higher metabolic capabilities thus affecting the final quality of the traditionally fermented product (Ayed *et al.*, 2002). Feng *et al.*, (2016) isolated lactobacilli from fermented vegetables and observed the same trend of *L. plantarum* being the dominant isolates. Liu *et al.*, also isolated lactic strains from pickled vegetables with unique probiotic characteristics.

### Resistance to simulated gastrointestinal juices

In order for probiotic candidates to exert their beneficial activity, they should survive exposure to gastrointestinal environment of low pH and other paramakers. The viable counts of all isolates of the different selected strains of *Lactobacillus* species for this test were less than or equal to 1 log CFU/ml as compared with the zero time count (7-8 log CFU/ml). This was noted at both pH 3 and 8 after 3 h and 4 h exposure, respectively. This high resistance was observed among *L. plantarum* 1 Y2, *L. pentosus* L 25 and *L. rhamnosus* Y1 isolates (Table 3). Some *L. plantarum* 1 namely L23, L24 were not as tolerant as others where at pH 4 viable counts were almost zero after 4h at pH 8. Maragkoudakis *et al.* (34.38) tested acid resistance of probiotic bacteria isolated from different sources and reported results in agreement with those reported herein. Abbas and Mahasneh isolated *Lactobacillus* isolates from camel milk and they had a similar trend in tolerance to gastrointestinal juices as those observed in this investigation. *L.plantarum* from fermented vegetables and it was able to show high survival rate after exposure to pH 2.5 and 8.0 for 3 and 4 hours respectively.

### Antagonistic activity against pathogens

All isolates inhibited the pathogenic target bacteria irrespective of being gram positive *B.cereus* or MRSA or gram negative *E.coli* or

*Salmonella typhimurium* with varying degrees (Table 4). This is associated with the ability of these isolates to produce bacteriocins and other antimicrobials (Messaoudi *et al.*, 2013) which needs further testing. Antibacterial substance production is a functional property to characterize probiotics (Shah, 2007). The ability to produce such compounds is very basic for competitive exclusion of pathogens and is a critical characteristic for probiotic bacterial candidates. Of interest is the superior antagonistic activity of *L. rhamnosus* Y1, *L. plantarum* 1 Y2, R21, R22, L23, L24 and *L. pentosus* L 25. It is notable that *L. rhamnosus* Y1, though greatly inhibited both *B.cereus* and *S.typhimurium*, it mildly inhibited MRSA. R22 *L.plantarum* 1 also highly inhibited *E.coli* and *B.cereus*.

### **Resistance to bile salts**

Most of *L.plantarum*1 isolates L23, L24 and Y2 were highly resistant to bile salts at the range of 0.3-2% after 24 h of exposure with little viable count reduction to the level of less than 1 log cycle (Table 5). However, *L. rhamnosus* Y1, lost about 50 percent of viability after exposure to 1 and 2% for 24 h. Since bile plays a role in the defenses of the gut, hence, bile tolerance is a paramount marker in choosing probiotic bacterial strains (Charteris *et al.*, 2000).

Sanders *et al.*, 1996 demonstrated the ability of lactobacilli to grow and metabolize at normal physiological bile concentrations of the gastrointestinal environment. Other reported the effect of food nature and components in the intestine in enhancing probiotics resistance to bile salts. Abbas and Mahasneh, 2015 reported variable degrees of survival of probiotic isolates among different *lactobacillus* species from camel milk. Studying fermented Korean Kimchi (2016) isolated *L.plantarum* strains surviving at high bile concentrations.

### **Antibiotic susceptibility**

Table 6 presents the minimum inhibitory concentration (MIC) breakpoints for the isolates. Strains were considered resistant if they had higher breakpoints compared with that of the European Food Safety Authority. Most tested isolates showed resistance to tetracycline, streptomycin, kanamycin and trimethoprim. *L.rhamnosus* Y1, the most sensitive it was resistant to 4 antibiotics and sensitive to 4 others. *L.plantarum* 1 M8 was sensitive to 3 antibiotics and resistant to 5 others. *L. pentosus* M5 and M9 strains were the most resistant isolates exhibiting resistance to 8 antibiotics of the 8 tested. The observed resistance in this study was in agreement with previous reports about *Lactobacillus* in general (Ashraf *et al.*, 2016). Tetracycline and streptomycin resistance among isolates was higher than that reported by others (Choi *et al.*, 2003) and in agreement with Ammor *et al.*, (Ammor *et al.*, 2008) reported high resistance among *Lactobacillus* isolates towards aminoglycosides (gentamicin and kanamycin) which was observed in this study. Resistance to such antibiotics is considered natural and intrinsic in lactobacilli due to it being chromosomally encoded. All *Lactobacillus* strains isolated from traditionally fermented cucumber and cabbage were resistant to streptomycin, vancomycin and gentamycin. Such resistance and in other studies are considered to be intrinsic and therefore not transmissible (Morrow *et al.*, 2012; Kumar *et al.*, 2012).

### **Adhesion ability of selected *Lactobacillus* isolates to Caco-2 cells**

The adhesion ability of probiotic lactobacilli to epithelial cells provides substantial benefits such as exclusion of pathogens. It also plays a role in the immunomodulation of the host (Haghshenas *et al.*, 2014) Adhesion to Caco-2 cells of isolates tested in this study varied between species and within species.

**Table.1** Some characteristic of bacterial isolates; all were non-spore forming rods

Isolate	Gram reaction	Catalase	Oxidase	Hemolysis	DNase	Gelatinase
Y1	+	-	-	-	--	
Y2	+	-	-	-	--	
R21	+	-	-	-	--	
R22	+	-	-	-	--	
L23	+	-	-	-	--	
L24	+	-	-	-	--	
L25	+	-	-	-	--	
M4	+	-	-	-	--	
M5	+	-	-	-	--	
M6	+	-	-	-	--	
M7	+	-	-	-	--	
M8	+	-	-	-	--	
M9	+	-	-	-	--	
M11	+	-	-	-	--	
B17	+	-	-	-	--	
B18	+	-	-	-	--	

**Table.2** Biochemical identification of *Lactobacillus* isolates according to Morphological and biochemical tests using API CH 50 strips

Isolate Code	Species Identification
Y1	<i>Lactobacillus rhamnosus</i>
Y2,R21,R22,L23,L24	<i>Lactobacillus plantarum</i> 1
L25,M4,M5	<i>Lactobacillus pentosus</i>
M6,M8	<i>Lactobacillus plantarum</i> 1
M11,B17	<i>Lactobacillus brevis</i> 3 and 1
B18	<i>Lactobacillus salivarius</i>

**Table.3** Effect of gastric and intestinal juices on viability of *Lactobacillus* isolates

Isolate	Zero count	3h count Ph3	4h count pH8
<b>Y1</b>	8.74 ± 0.14	8.01 ± 0.07	7.05 ± 0.13
<b>Y2</b>	8.35 ± 0.06	7.87 ± 0.04	7.33 ± 0.07
<b>R21</b>	8.23 ± 0.03	6.10 ± 0.17	0.00 ± 0.00
<b>R22</b>	8.50 ± 0.02	7.47 ± 0.05	6.52 ± 0.07
<b>L23</b>	7.53 ± 0.21	6.30 ± 0.30	0.00 ± 0.00
<b>L24</b>	9.56 ± 0.02	6.79 ± 0.79	0.00 ± 0.00
<b>L25</b>	9.46 ± 0.10	6.49 ± 0.20	6.00 ± 0.00
<b>M11</b>	7.80 ± 0.10	7.40 ± 0.22	6.20 ± 0.06
<b><i>L. reuteri</i></b>	7.10 ± 0.17	7.52 ± 0.09	7.06 ± 0.25
<b>M4 – M9</b>	Not tested	Not tested	Not tested
<b>B7-B18</b>	Not tested	Not tested	Not tested

Results are presented as log<sub>10</sub> CFU/ml + S.D,N=2 after 3h and 4h exposure at pH 3 and 8 respectively



**Table.4** Antagonistic activity of some *Lactobacillus* isolates against pathogenic bacteria as Indicator

Inhibition zone diameters (mm) of indicator strains				
Bacterial isolate	<i>B. cereus</i>	<i>E. coli</i>	MRSA	<i>S. typhimurium</i>
Y1	60 ± 0.0	45 ± 7.1	15 ± 0.0	50 ± 0.0
Y2	40 ± 0.0	36 ± 0.0	30 ± 0.0	46 ± 0.0
R21	43 ± 1.4	50 ± 0.0	50 ± 0.0	38 ± 2.8
R22	50 ± 0.0	64 ± 0.0	35 ± 7.1	40 ± 0.0
L23	50 ± 0.0	44 ± 2.8	14 ± 0.0	30 ± 0.0
L24	42 ± 0.0	40.5 ± 2.1	40 ± 0.0	50 ± 0.0
L25	50 ± 0.0	38 ± 2.8	20 ± 0.0	28 ± 0.0

Inhibition zone diameters (mm) are presented as mean + S.D.; n=2

**Table.5** Tolerance of some selected *Lactobacillus* isolates to varying concentrations of exposure and bile salts after 24 h of anaerobic incubation

Isolate	Bile concentration%							
	0.3		0.5		1.0		2.0	
	0 h	24 h	0 h	24 h	0 h	24 h	0 h	24 h
Y1	6.59 ± 0.11	8.80 ± 0.30	6.04 ± 0.03	6.39 ± 0.07	5.10 ± 0.07	4.52 ± 0.03	5.47 ± 0.04	3.77 ± 0.09
Y2	6.07 ± 0.45	5.60 ± 0.00	6.30 ± 0.17	5.18 ± 0.03	5.90 ± 0.21	5.18 ± 0.05	5.94 ± 0.30	5.18 ± 0.08
R21	6.33 ± 0.07	7.33 ± 0.35	5.92 ± 0.03	7.67 ± 0.01	5.10 ± 0.17	4.95 ± 0.00	5.00 ± 0.00	4.20 ± 0.17
R22	5.73 ± 0.23	5.38 ± 0.03	5.78 ± 0.15	3.58 ± 0.51	6.06 ± 0.11	3.59 ± 0.06	5.76 ± 0.14	3.16 ± 0.04
L23	6.23 ± 0.07	8.43 ± 0.23	6.17 ± 0.13	7.77 ± 0.28	6.50 ± 0.16	7.34 ± 0.03	6.16 ± 0.02	6.38 ± 0.00
L24	6.47 ± 0.28	9.00 ± 0.00	6.18 ± 0.14	8.15 ± 0.04	6.14 ± 0.10	6.32 ± 0.10	6.12 ± 0.26	5.26 ± 0.00
L25	6.44 ± 0.06	8.40 ± 0.36	6.40 ± 0.02	7.41 ± 0.09	6.32 ± 0.19	7.20 ± 0.00	6.28 ± 0.11	5.70 ± 0.00

Results are presented as mean + S.D. of viable counts at zero and 24 h exposure to bile, n = 2

**Table.6** Antibiotic susceptibility profiles of some *Lactobacillus* isolates of probiotic potential.

Antibiotic breakpoint <sup>a</sup> (µg/ml)								
Isolate	A (2)	C (4)	E (4)	G (1)	K (32)	S (16)	Te (16)	Tr (16)
Y1	S	R	S	S	R	R	R	S
M4	S	S	R	R	R	R	R	S
M5	R	R	R	R	R	R	R	R
M6	R	S	R	R	R	R	R	R
M7	R	R	R	S	S	R	R	R
M8	S	R	S	S	R	R	R	R
M9	R	R	R	R	R	R	R	R
<i>L. reuteri</i> DSMZ 20056	S	R	S	S	S	S	R	S

<sup>a</sup>The breakpoints for *Lactobacillus* sp. by SCAN category. Minimum Inhibitory Concentration (MIC) equal to or higher than the breakpoint is considered as resistant. (S): Susceptible; (R): Resistant; (A): Ampicillin; (C): Ciprofloxacin; (E): Erythromycin; (G): Gentamicin; (K): Kanamycin; (S): Streptomycin; (Te): Tetracycline and (Tr): Trimethoprim

**Table.7** Relative adhesion ability to Caco-2 cells of some selected *Lactobacillus* isolates.

Test strain	Log CFU/ml Adhered	Adhesion %
<i>E.coli</i>	6.39 +_ 0.17	100%
<i>L.reuteri</i>	4.83+_ 0.93	75 %
M 5	6.40 +_ 0.13	100%
M 6	5.30 +_ 0.28	83%
M 8	3.26 +_ 0.24	51%
M 11	5.22 +_ 0.36	82 %
B 17	3.88 +_ 0.30	61%
B 18	3.42 +_ 0.26	54 %

*Escherichia coli* and *Lactobacillus reuteri* DSMZ 20056 were included as a pathogen ( 100% relative adhesion) and known probiotic strains respectively

Considering *E.coli* relative adhesion in our test as 100 percent, *L. pentosus* M5, *L. plantarum* M6 and *L. brevis* M11, all showed relative adhesion ability above 80% (Table 7) and it was higher than *L. reuteri* DSMZ 20056 which is a probiotic strain. Low adhesion values were recorded for M8 (51%) and *L. salivarius* B18 (54%).

Studying *Lactobacillus* strains isolated from fermented vegetables recorded much lower adhesion values ranging from 0.45 - 12.27 %

in most cases, it is believed that variations are strain specific and is linked to differences in cell surface structures (Garcia-Ruiz *et al.*, 2014; Hu *et al.*, 2015). Finally, taking all of the above results in consideration, it is clear that traditionally fermented FOODS and vegetables and Makdoos in this case presents a promising source of improved probiotic lactobacilli due to stresses exerted upon the bacteria during the course of ripening of the mix of fermented vegetables. Tested strains isolated in this study are positively rated as

suitable probiotics, however additional *in vitro* studies would substantiate fully their medicinal and otherwise uses.

In conclusion, it is becoming increasingly obvious that the traditional fermented foods offer unlimited reservoir of beneficial probiotics acting upon several targets to improve and protect human health. However, *in vitro* results should be substantiated by *in vivo* studies

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