



Screening, Identification and Production of L-glutaminase by *Lactobacillus reutri* Isolated from feces of healthy adults and effect it on *P.aerogenosa* wound infection

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ABSTRACT

Lactobacillus reutri may be employed as a functional antibacterial agent. Optimizing the antibacterial activity of L-glutaminase extracted from *L. reutri* led to an increase in the enzyme's synthesis in low-medium production media. By employing *L. reutri* as antibiotics and antibacterial biofilms, this work aims to increase the production of high-activity L-glutaminase that has been isolated from the bacterium. In ion exchange chromatography, the specific activity was 4.7 U/mg of protein, with a yield of 65% and 6.6 times purification. In gel filtration, the specific activity was 10 U/mg of protein, with a yield of 60% and 16.6 times purification and the molecular weight of glutaminase was found to be 34-40 KD determined by sodium dodecyl sulfate polyacrylamide. The results indicated that L-glutaminase that was isolated from *L. reutri* had strong antibacterial and antibiofilm properties. The antibacterial activity of isolated L-glutaminase was shown to be substantial ($P < 0.05$) when applied to *P. aeruginosa* isolates. Furthermore, the antibiofilm activity of purified L-glutaminase was found to be more potent than that of piperacillin tazobactam, the reference standard drug ($P < 0.05$). This finding may prove promising for the development of natural probiotics or other treatments for bacterial illnesses in the future.

Key word: Lactic acid bacteria (LAB), Biofilms, Antibiotic, Antibiotic Susceptibility. Probiotics, L- Glutaminase.

INTRODUCTION

According to Charteris *et al.* (1998), probiotics are living microorganisms with nutritional and physiological benefits. One hundred years ago, Elie Metchnikoff was the first person to do research on probiotics (Muhammad and Ahmad, 2018). Many probiotic microorganisms are composed of lactic acid bacteria (LAB), including *Bifidobacterium* sp., *Lactobacillus* sp., *Streptococcus* sp., *Enterococcus* sp., and *Pediococcus acidilactici* (Sharma *et al.*, 2014). L-glutaminase (EC3.5.1.2; L-glutaminase amid hydrolase). L-glutaminase breaks down L-glutamine into an ammonia group and glutamic acid (Sarkar *et al.*, 2014; Ayodeji *et al.*, 2019). Glutamine amino transferases, often known as glutamines, catalyze the bacterial conversion of glutamine to glutamate. Acid resistance and the production of the bioactive metabolites glutamate, -amino butyrate (GABA), and glutamyl γ gammapeptides are both influenced by glutamine deamination. Both prokaryotes and eukaryotes rely heavily on L-glutaminase for their nitrogen metabolism (Katikala *et al.*, 2009). Potential sources of L glutaminase include bacteria, fungi, and yeast. It is preferable to use these microorganisms for L-glutaminase synthesis due to their easy processing, low production costs, and straightforward growth requirements (Sarkar *et al.*, 2014). L-glutaminase is a very significant medicinal substance that is employed to treat illnesses like cancer and to enhance other organ ailments like gastrointestinal issues and

MATERIAL AND METHODS

Isolation and identification of *L. reuteri* from feces

Fifty six isolates of *Lactobacillus spp* obtained from 111 samples of feces of healthy adults volunteer (22 to 30 years) The biochemical reactions of the *L. reuteri* isolates were, as displayed in table 3-3. All the bacterial isolates

antibacterial effects. Numerous earlier investigations used bacteria from various species and environments, including *Lactobacillus ferment* and *Aspergillus niger*, to screen, isolate, and purify L glutaminase. (Woraharn *et al.*, 2014; Sushma *et al.*, 2017; Dutt *et al.*, 2014). The genus *Lactobacillus* is a member of the big LAB group and is frequently utilized in food as probiotics and as starters for fermented foods (Arena *et al.*, 2016). Among the lactobacilli, *Lactobacillus salivarius* is a type of Gram-positive, anaerobic rods that do not form spores. It is commonly found in human saliva and other oral secretions, as well as in the digestive tracts of both humans and animals (Caralampopoulos *et al.*, 2009). It's thought that many LAB have probiotic properties. "Live microorganisms which, when administered in adequate amounts, confer a health benefit on the host" is the definition of probiotics (Guarner *et al.*, 1998). Numerous LAB-produced antimicrobial substances, such as bacteriocins, organic acids, diacetyl, and hydrogen peroxide, are capable of inhibiting a wide range of harmful microbes (Bamidele *et al.*, 2013). A commensal bacteria called *L. gasseri* is present in the female lower urogenital tract and the human GI tract (Walter, 2008; Matsuda *et al.*, 2018). The effect on women's health has been studied recently (Matsuda *et al.*, 2018; Phukan *et al.*, 2018). Mahdi *et al.* (2024) conducted a study on the effectiveness of l-glutaminase derived from clinical isolates of *L. gasseri* BRLHM as an antibacterial agent against *P.aeruginosa* that is resistant to many drugs.

were oxidase, catalase, gelatinase and citrate negative; *Lactobacillus* forms a clear zone on MRS medium that is supported with CaCO_3 , because of their production of lactic acid that would combine with calcium then precipitate. All

Lactobacillus isolates were capable to grow in the existence of 2.5 and 5 % NaCl while at 6.5 % NaCl they were not capable to grow. According to the morphological and biochemical tests, the isolates were identified as Lactobacillus spp. The slides

Preparation of crude L-glutaminase BRLG:

L.gasseri in its raw form With a few modifications, L-glutaminase BRLG was produced in accordance with the methodology described by Padma *et al.* (2009). Subsequently, the culture was grown in L-glutamine broth (made by dissolving 1 g yeast extract, 5 g glucose, 5 g glutamine, 6 g Na₂HPO₄.2H₂O, 3 g KH₂PO₄, 0.49 g MgSO₄, 0.05 g NaCl, 0.002 g CaCl₂, 20 gm of agar, and 2.5% w/v ethanolic phenol red solution 0.06 ml in 1 liter of deionized distilled water. The pH was then adjusted to 6.2 and the broth was autoclaved at 121 c for 10 minutes.) at 37°C for 24 hours under anaerobic conditions. The cells were separated by centrifugation at 8000 rpm for 15 minutes at 4°C.

RESULTS

Isolation and identification of *L. reuteri* from feces

Fifty six isolates of *Lactobacillus spp* obtained from 111 samples of feces of healthy adults volunteer (22 to 30 years) The biochemical reactions of the *L. reuteri* isolates were, as displayed in table 1. According to the

underwent heat fixation, air drying, and Gram staining. The slides were next examined in accordance with , which studies the size, shape, color, and edge of colonies grown on MRS agar (Atlas *et al.* ,1995; Goldman and Green ,2015).

Aliquots of the supernatant were then pH-corrected to 6.2 and kept at 4°C until analysis. Semi-quantitative screening was conducted using the Gulati *et al.* (1997) approach. The procedure outlined by Mannan *et al.* (1995) was followed in the quantitative assay to ascertain the enzyme activity in every isolate. Imada *et al.*'s (1973) assay of enzyme activity was determined utilizing these procedures.

Determination of optimum condition for L-glutaminase production:

The optimum of carbon source, nitrogen source, pH, and temperature and incubation period was estimated according to the method described by Savitha *et al* (2016).

morphological and biochemical tests, the isolates were identified as Lactobacillus spp. Dependent on the results of API 50 KIT six species were identified as the following L. plantarum 22 isolates (39%) L. fermentum 13 isolates (23%), L. reuteri 12 isolates (21%), L. rhamnosus 6 isolates (10%) L. paracasei 2 isolates (3%) and L.gasseri 1 (1%) as shown in fig 1.

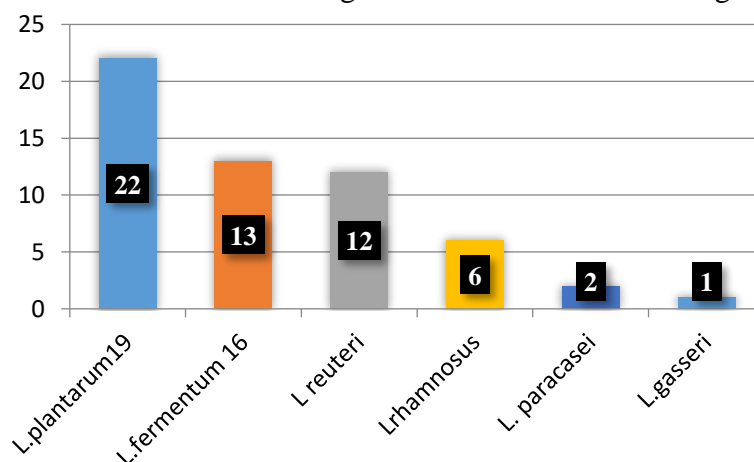


Fig (1): Percentage of Lactobacillus spp. isolated from feces of healthy adults volunteer

Semi Quantitative Screening of L- glutaminase produced from *L. reuteri* isolates isolated from feces

By using the L-Glutamine agar media containing phenol red were used for assay Semi Quantitative Screening of L- glutaminase produced from *L. reuteri* isolates isolated from feces . Results

indicated a correlation between the pink color and the amount of generated enzyme. *L. reuteri* isolate number 3 could be regarded as the highest L- Glutaminase producer because it had the largest a visible pink zone when compared to the other isolates. Result showed that the isolates were L- glutaminase producers, as shown in (Table 1).

(Table 1).with different levels of productivity.

Isolates	Pink zone (Mean \pm SD) (mm)
<i>L. reuteri</i> no.1	10.22 \pm 1.93
<i>L. reuteri</i> no. 2	7.49 \pm 1.64
<i>L. reuteri</i> no. 3	13.57 \pm 3.8
<i>L. reuteri</i> no.4	0 \pm 0
<i>L. reuteri</i> no. 5	11.42 \pm 2.07
<i>L. reuteri</i> no. 6	8.87 \pm 0.56
<i>L. reuteri</i> no.7	0 \pm 0
<i>L. reuteri</i> no. 8	9.89 \pm 3.11
<i>L. reuteri</i> no. 9	12.39 \pm 2.08
<i>L. reuteri</i> no.10	11.52 \pm 2.23
<i>L. reuteri</i> no. 11	6.71 \pm 1.22
<i>L. reuteri</i> no. 12	13.03 \pm 1.66
<i>L. reuteri</i> no. 12	13.03 \pm 1.66

Quantitative Screening of L- glutaminase produced from *L. reuteri* isolates

After Semi Quantitative screening, the specific L- glutaminase activity for *L. reuteri* isolates in liquid media were identified. The isolate No.3 provided

the highest (89.06 U/mg protein) Specific activity. Moreover, isolate No. 8 showed the lowest specific activity 33.98 U/mg proteins. According to these results, the isolate No. 3 was selected to be used for improving its ability in L-glutaminase production .

Isolates	Enzyme activity	Protein mg/n	
			Specific activity (U/mg)
no.1	9.97 \pm 0.58b	0.122 \pm 0.02	78.66 \pm 3.42c
no. 2	5.32 \pm 2.42d	0.136 \pm 0.03	47.42 \pm 2.69d
no. 3	13.88 \pm 1.71a	0.161 \pm 0.01	89.06 \pm 1.99a
no.5	10.29 \pm 1.06b	0.143 \pm 0.01	81.05 \pm 3.06b
8	7.01 \pm 1.45c	0.122 \pm 0.01	33.98 \pm 2.14d

no.9	10.09±2.16ab	0.108±0.04	81.22±1.92b
no. 10	10.33±1.62b	0.139±0.01	79.55±0.93b
no.12	11.93±2.51ab	0.142±0.02	83.69±2.34a
LSD	5.439**	NS	6.38**

(Table 2).LSD test was used to calculate the significant differences between tested mean, the letters (A, B,C and D) LSD for rows represented the levels of significant, highly significant start from the letter (A) and decreasing with the last one. Similar letters mean there are no significant differences between tested mean

Extraction of L- glutaminase produced from *L. reuteri* isolate no.3

The *L. reuteri* isolate no. 3 was selected to be used for improving its ability in L-Glutaminase production by optimization. This approach is used to screen *P. fluorescens* for L-glutaminase synthesis. Variations in the genes responsible for producing this enzyme may account for specimen discrepancies in their ability to make it (Kumari *et al.*, 2015).*A. faecalis* is used to screen for L-glutaminase (Pandian *et al.*, 2014).Ion exchange chromatography (DEAE-Cellulose)

The results in figure 3-6 showed that their Ionic exchange chromatography patterns showed one protein peak in wash elution and one peak in stepwise elution. Only one peak among the stepwise elution peaks represented glutaminase activity appeared by measuring the absorbance at 280 nm of each fraction, at elution fractions tube glutaminase s (56, 1.3) with 0.6 M of NaCl.The fractions were pooled and tested for glutaminase activity with 0.65 U/ml, specific activity 4 U/mg a fold purification of 6.6 and protein yield of 65 % as shown in table 3-5.LGlutaminase from *Bacillus cereus* were purified using a DEAE-cellulose exchanger.

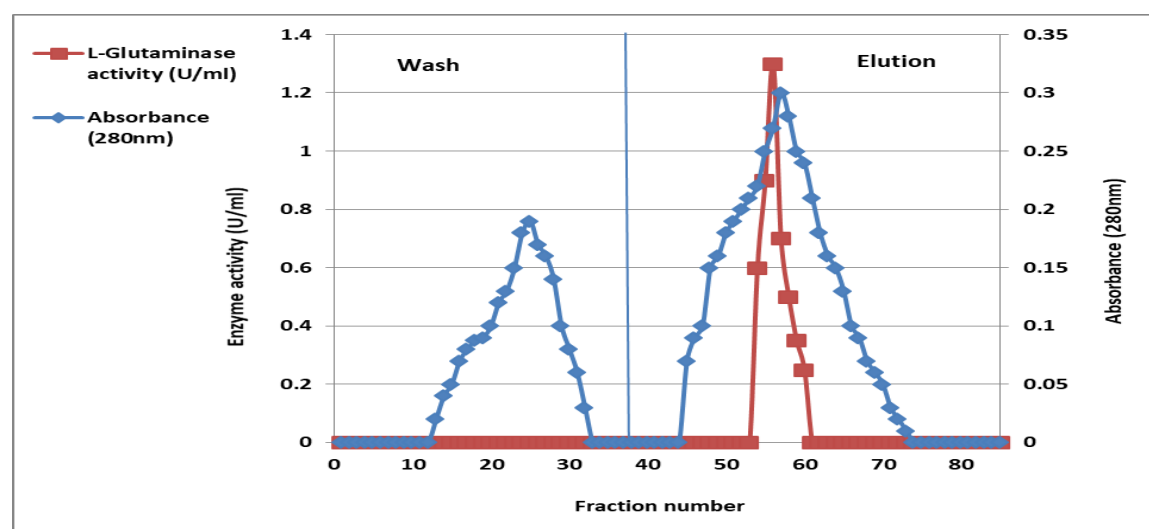


Fig (2): Ionic exchange chromatography for Rhamnocin LHM through DEAE cellulose column (2x 25) cm. The column was calibrated with 0.5 M phosphate buffer, pH7, flow rate 60ml/hrs and 3 ml fraction, eluted with stepwise (0.1-1) M NaCl.

Gel filtration chromatography

Further purification carried out by a gel filtration using Sephadex G-100 column L- glutaminase active fractions from DEAE-cellulose were pooled and passed through gel filtration column. The fractionation yielded one protein peak as absorbance reading at 280nm, this peak series

(19,1) contained glutaminase activity **0.7 U/ml**, specific activity **10 U/mg** and the purification fold was **16.6** with yield of protein **60 %** as mentioned in figure 3-7 and table 3-5. Gel filtration is unusual in that fractionation is based on the relative sizes of protein molecules and is carried out utilizing porous beads as the chromatographic support.

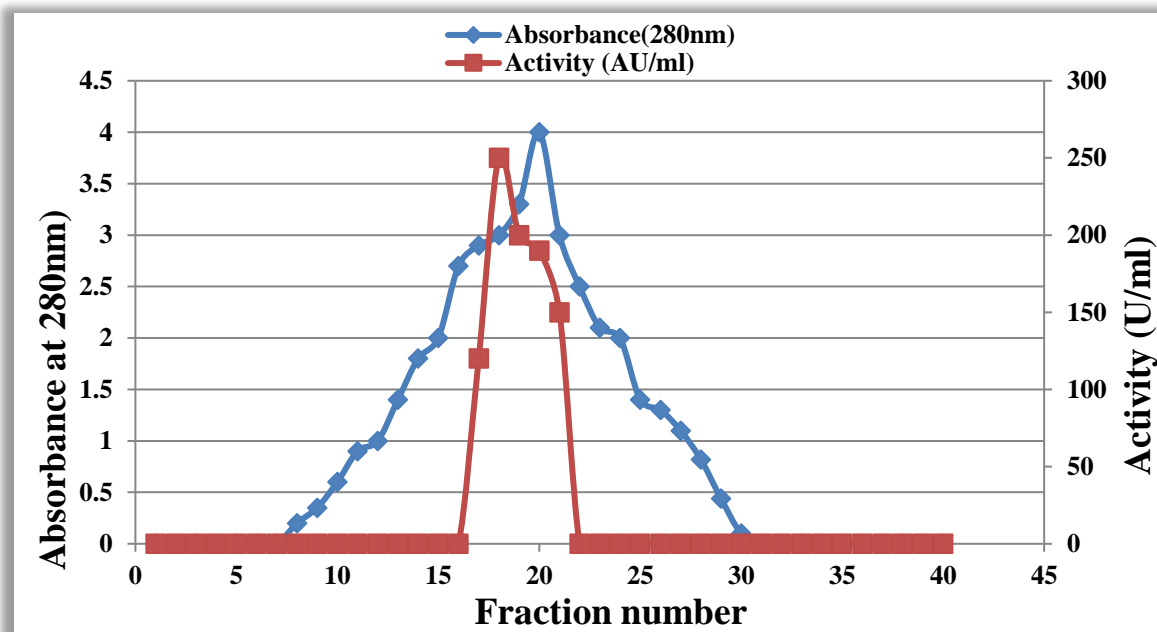


Fig (3): Gel filtration chromatography for purified l-glutaminase by using Sephadex G- 100 column (2x35) cm. The column was calibrated with 0.5 M phosphate buffer, pH7; flow rate 30 ml/hrs and 5 ml/fraction.

Steps of purification	Volume (ml)	Enzyme activity (U/ml)	Protein concentration (mg/ml)	Specific activity (U/mg)	Total activity (U)	Purification (folds)	Yield (%)
Crude enzyme	70	0.3	0.5	0.6	21	1	100
Ammonium sulphate precipitation 70%	14	1	0.45	2	14	3.3	66
DEAE-cellulose	21	0.65	0.15	4	13.6	6.6	65
Sephadex- G100	18	0.7	0.1	10	12.6	16.6	60

(Table 3).Equations for purification table

Determination of Molecular Weight of glutaminase

Sodium Dodecyl Sulfate-Polyacrylamide was used to determine the molecular weight of glutaminase. Gel electrophoresis by using SDS-PAGE, the

molecular weight of glutaminase was calculated using the standard curve that depicts the relationship between log of the molecular weight and relative mobility (Rm) for standard proteins. The molecular weight of glutaminase was found to 34-40 KD, as illustrated in fig (4 ,5)

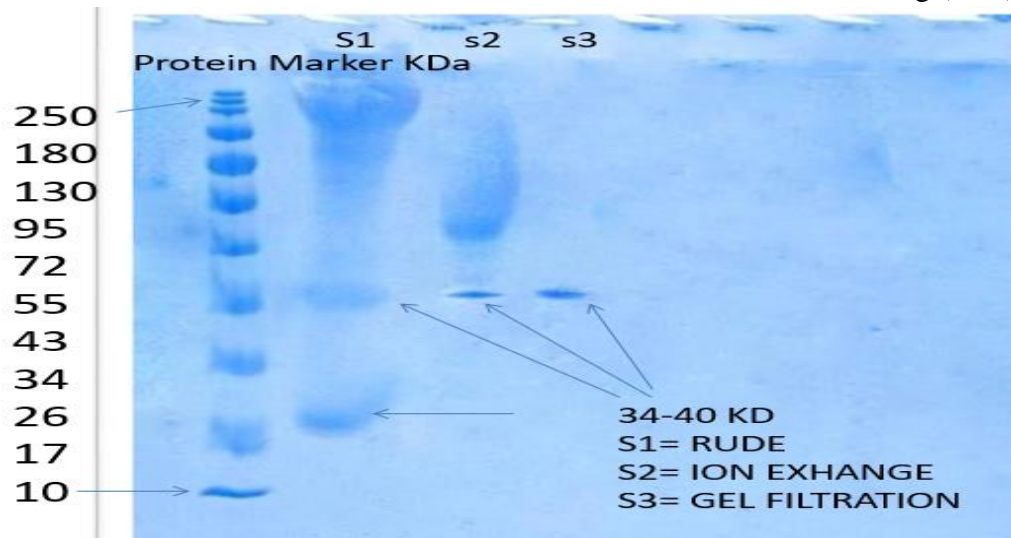


Figure (4): Sodium dodecyl sulfate-poly acrylamide gel electrophoresis (SDS-PAGE) of the purified Rhamnocin LHM.

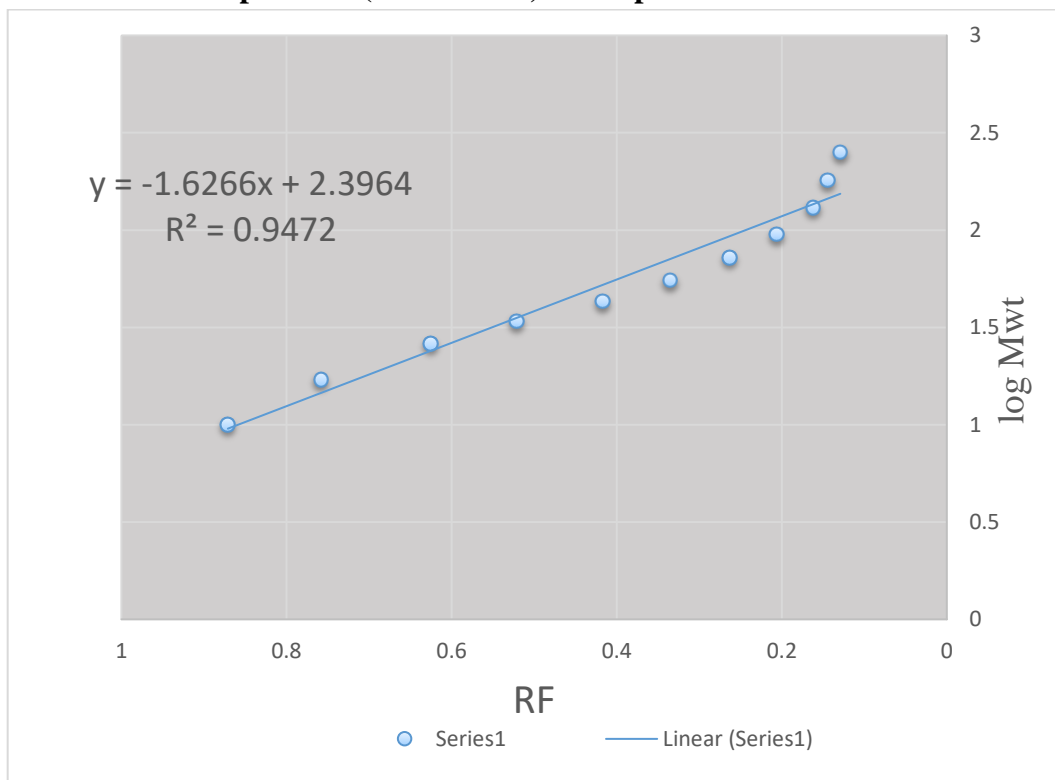


Fig (5): Determination of molecular weight of purified glutaminase.

Log Mwt: logarithm of molecular weight, Da: Dalton.

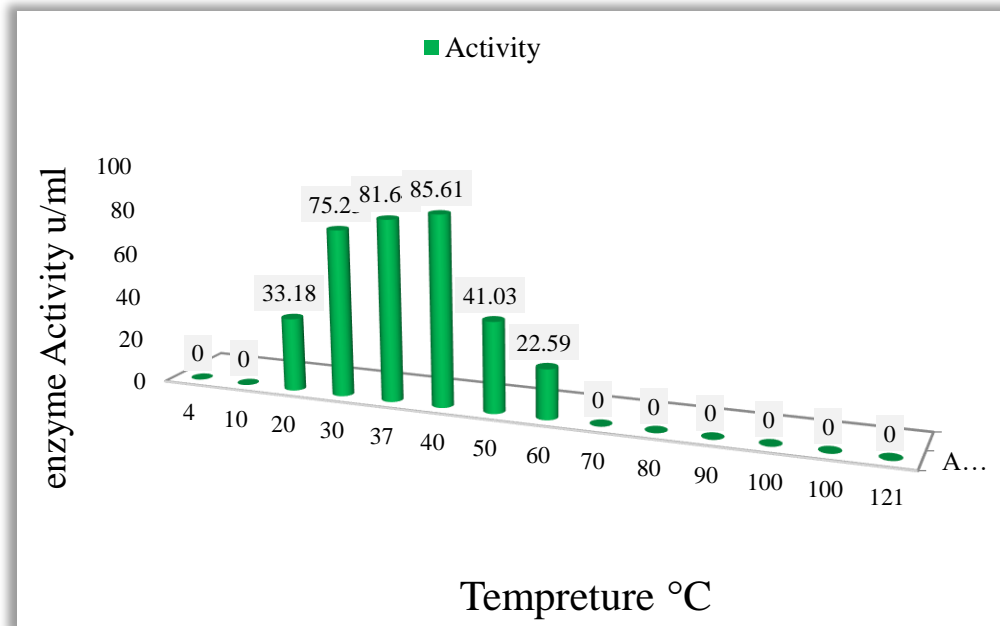


Fig (6): activity of purified L-glutaminase purified from L. reuteri isolate no.3 at different temperatures

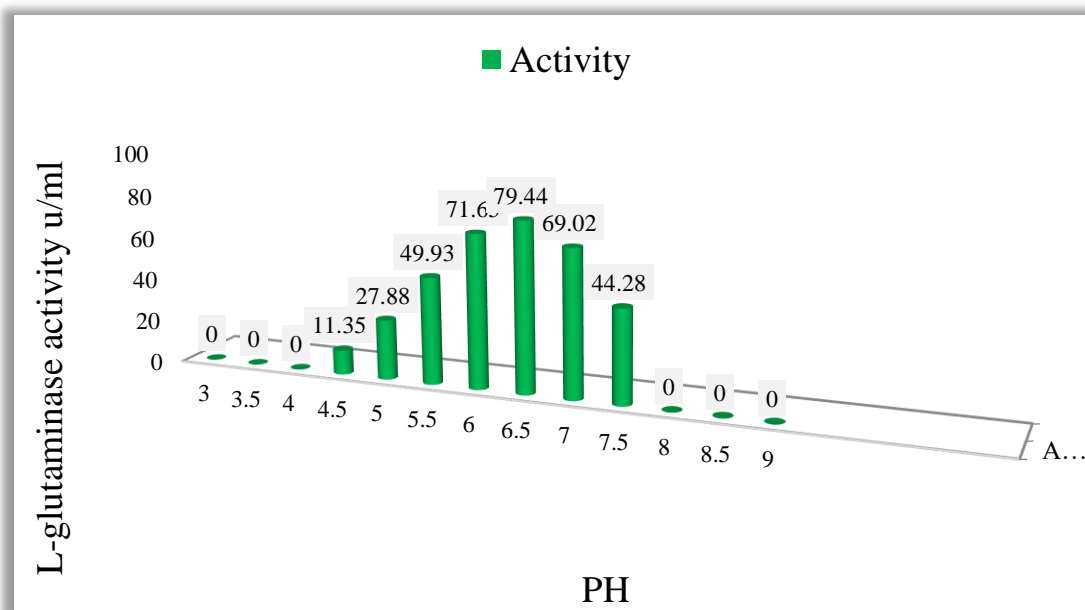


Fig (7) :- activity of L-glutaminase purified from L. reuteri isolate no.3 at different pH values.

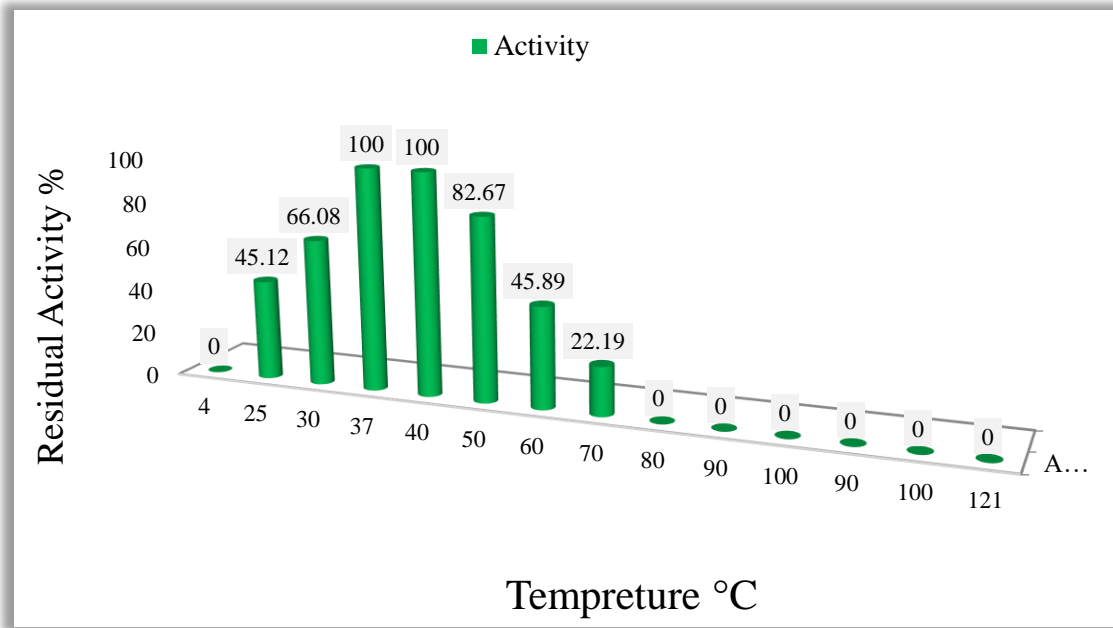


Figure (8): thermal stability of L-glutaminase purified from *L. reuteri* isolate no.3 at different temperatures

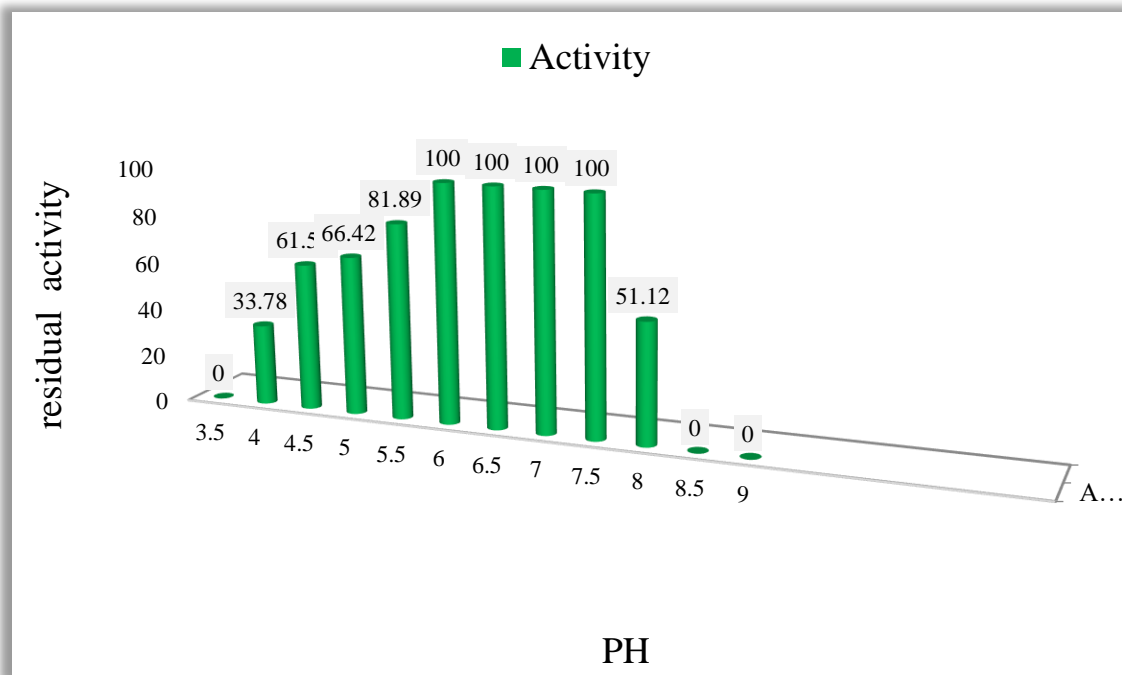


Fig (9): pH stability of L-glutaminase purified from *L. reuteri* isolate no.3 at different pH values.

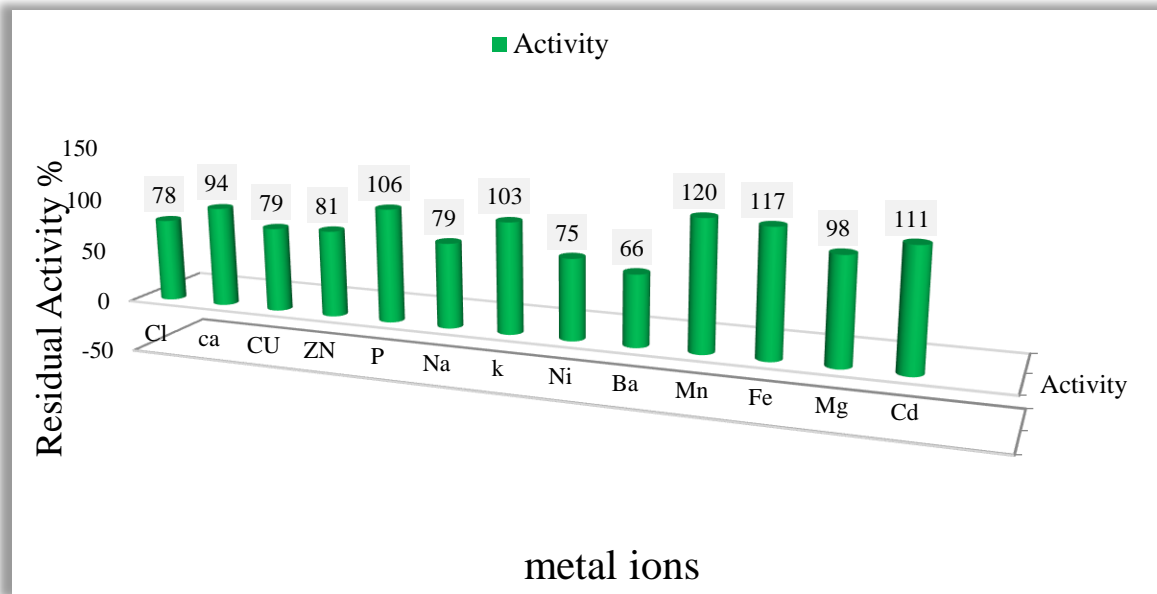


Fig (10): effect of metal ions on L-glutaminase purified from L. reuteri isolate no.3activity

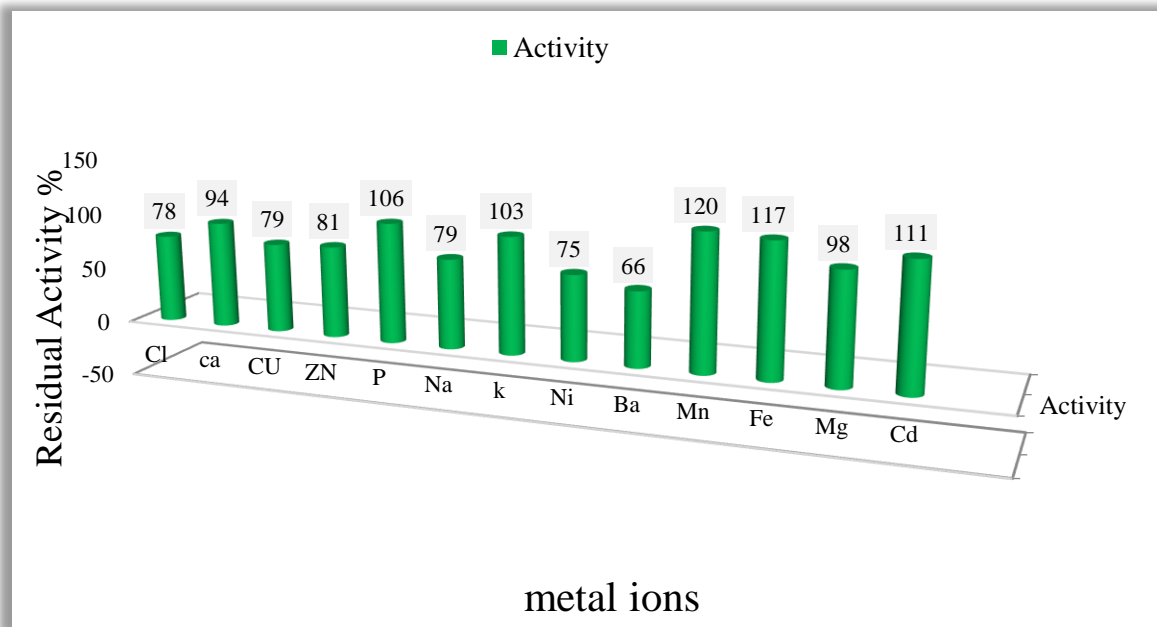


Fig (11): effect of metal ions on L-glutaminase purified from L. reuteri isolate no.3activity

Table (4): Antibacterial activity of L-glutaminase purified from *L reuteri* on *P. aeruginosa* isolates in-vitro

Isolates number	Concentration $\mu\text{g/ml}$				P value
	Crude L-glutaminase 64	Crude L-glutaminase 32	Purified L-glutaminase		
			64	32	
<i>P. aeruginosa 5</i>	C 12.07 \pm 2.11	C 14 .23 \pm 1.06	A 34.7 \pm 2.98	B 22.55 \pm 3.04	0.01 *
<i>P. aeruginosa9</i>	C 9.66 \pm 1.71	C 13.59 \pm 2.44	A 29.09 \pm 3.07	B 24.3 \pm 1.09	0.01 *
<i>P. aeruginosa24</i>	C 11.69 \pm 1.02	C 14.08 \pm 0.08	A 31.22 \pm 1.33	B 20.15 \pm 2.03	0.01 *
<i>P. aeruginosa31</i>	C 8.11 \pm 0.9	C 10.22.6 \pm 0.89	A 25.71 \pm 2.09	B 17.49 \pm 0.03	0.01 *
<i>P. aeruginosa33</i>	C 13.01 \pm 1.92	C 15.02 \pm 0.08	A 30.51 \pm 0.7	B 19.12 \pm 1.76	0.01 *
<i>P. aeruginosa58</i>	C 7.05 \pm 0.9	C 10.38 \pm 0.6	A 24.06 \pm 1.05	B 20.11 \pm 0.9	0.01 *
P value	NS	NS	NS	NS	

t test was used to calculate the significant differences between tested mean, (*: significant differences)

Letters (A, B and C) LSD for rows represented the levels of significant, highly significant start from the letter (A) and decreasing with the last one. Similar letters mean there are no significant differences between tested mean.)

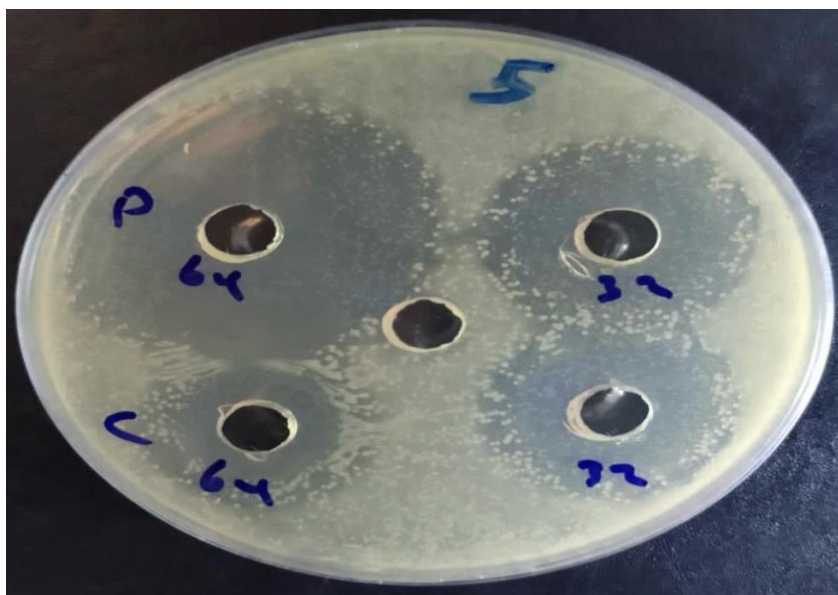


Figure (12): Antibacterial activity of crude and purified L-glutaminase against *P. aeruginosa*. A: at concentration 64 $\mu\text{g/ml}$, B: at concentration 32 $\mu\text{g/ml}$.

Determination of MIC and Sub-MIC of Piperacillin tazobactam and purified L-glutaminase

The six *P. aeruginosa* isolates were selected according to it is the ability to greater biofilm

production and it is the capability of greater antibiotics resistance. (*P. aeruginosa* no. (5,9,24,31,33,58) isolates. Piperacillin tazobactam and purified L-glutaminase were tested by determining the MIC utilizing a microtiter plate with Resazurin stain.

Table (5): MIC and sub MIC of Piperacillin tazobactam and purified L-glutaminase against *P. aeruginosa* isolates.

Treatment $\mu\text{g/ml}$	Ps. 5	Ps. 9	Ps. 24	Ps. 31	Ps. 33	Ps. 58
MIC of Piperacillin tazobactam	32	256	64	64	16	8
Sub-MIC of Piperacillin tazobactam	16	128	32	32	8	4
MIC of purified L-glutaminase	8	64	32	32	128	256
Sub-MIC of purified L-glutaminase	4	32	16	16	64	128

Inhibitory effect of purified L-glutaminase & Piperacillin tazobactam on biofilm formation by *P. aeruginosa* isolates.

In this study, the high resistance to antibiotics and high producing biofilm clinical isolates were chosen (5,9,24,31,33, and 58). To evaluate the antibiofilm activity of purified L-glutaminase from purified and Piperacillin tazobactam, sub MIC were used. Optical density exhibited significant differences before and after treatment. ($P < 0.05$) as displayed in table 5 purified L-glutaminase activity as antibiofilm agent was statistically significant higher than the activity of the

Piperacillin tazobactam ($P < 0.05$) in vitro. On the other hand, after comparing among isolates, the results revealed that the isolates *P. aeruginosa* no.5, no.9 and no.58 were the most biofilm producing isolates after comparing with other clinical isolates.

Table (6): Effect sub-MIC of purified L-glutaminase and Piperacillin tazobactam on biofilm formation from *P. aeruginosa*

No. of Isolates	O. D. Before treatment	O. D. after Treatment with sub MIC		P value
		Piperacillin tazobactam	purified L-glutaminase	
Ps. 5	1.622 \pm 0.01 A	0.71 \pm 0.08 C	0.47 \pm 0.17 D	SIG 0.05
Ps. 9	1.738 \pm 0.22 A	0.53 \pm 0.05 C	0.34 \pm 0.05 E	SIG 0.05
Ps. 24	1.691 \pm 0.19A	0.59 \pm 0.11C	0.29 \pm 0.08E	SIG 0.05

Ps.31	1.42±0.46A	0.61±0.25 C	0.48±0.09D	SIG 0.05
Ps.33	1.29±0.81 B	0.39±0.09 D	0.29±0.07 E	SIG 0.05
Ps.58	1.622±0.92A	0.55±0.12 C	0.41±0.12 D	SIG 0.05
P value	0.05	0.05	0.05	-----

ANOVA test was used to calculate the significant differences between tested mean, the letters (A, B,C,D and E) for rows represented the levels of significant, highly significant start from the letter (A) and decrease through the last one. Similar letters mean there are no significant differences between tested mean. (not significantly)

CONCLUSION

Lactobacillus reutri are good probiotics and a good antibiotic for pathogenic bacteria, especially *P. aeruginosa*, as well as an anti-biofilm, and It is a good producer of the L-glutaminase enzyme, which allows the possibility of using them as a treatment in the future.

DISCUSSION:

According to the measurement of L-Glutaminase activity in cell-free extracts, 5 Unit/mg of *L. gasseri*-specific activity and L-Glutaminase were produced. In applied and industrial microbiology, lactobacillus strains remain the most common bacteria that produce enzymes; these organisms represent a significant source of extracellular industrial enzymes (Ziadé *et al.*, 2003). Since glutamine amino acid was the only carbon and nitrogen source in glutamine agar, organisms that could produce glutaminase could thrive on this medium. The PH indicator changed from yellow to pink in color. Around colonies, a pink zone was observed due to the activity of the L glutaminase enzyme, which broke the L glutamine amino acid and released ammonia, raising the pH level (Sarkar *et al.*, 2014). Using this method to screen for L-glutaminase synthesis from *P. fluorescens*, the differences in specimens' abilities to make this enzyme could be attributable to genetic variants in

the genes responsible for producing this enzyme (Kumari *et al.*, 2015). Screened L-glutaminase from *A. faecalis*. Many probiotic microorganisms are composed of lactic acid bacteria (LAB), including *Bifidobacterium* sp., *Lactobacillus* sp., *Strep. thermophilus*, *Enterococcus* sp., and *Pediococcus acidilactici* (Pandian *et al.*, 2014; Sharma *et al.*, 2014). High-level pure *L. gasseri* BRLHM glutaminase was shown to have antioxidant properties (Hasson *et al.*, 2021). Genetic variations in the genes that create this enzyme may be the cause of the variances in each specimen's capacity to manufacture it (Kumari *et al.*, 2015). This technique is employed to screen *P. fluorescens* for the synthesis of L-glutaminase. *A. faecalis* is used to screen for L-glutaminase (Pandian *et al.*, 2014). According to Singh *et al.* (2016), purified L-glutaminase exhibited a specific activity of 7.44 U/mg and a purification fold of 14.8. Using ion-exchange chromatography, L-glutaminase was also extracted with a purification fold of 46 from *Streptomyces* sp. (Desai *et al.*,

2016). The enzymes produced by *Lactobacillus gasserii* were further purified and characterized using a chromatography method (Hasson *et al.*, 2021). Using gel filtration, L-glutaminase from marine *Brevundimonas diminuta* was isolated, achieving an activity of 60.15 U/mg and an enzyme recovery of 48.12% (Jayabalan *et al.*, 2010). The molecular weights of microbial L-glutaminase ranged between 35 ± 1 and 71 KDa, and electrophoresis is thought of as an additional purification step (Desai *et al.*, 2016; Savitha *et al.*, 2016). A few larger molecular mass exceptions have been found, including 50 KDa from *Streptomyces* sp. isolated from soil (Desai *et al.*, 2016; Savitha *et al.*, 2016); 64 KDa L-glutaminase from *Bacillus subtilis* was also reported (Jambulingam and Sudhakar, 2019). Whether it was crude or purified, the data indicate that a concentration of 64 $\mu\text{g/ml}$ was more effective against *P. aeruginosa* than 32 $\mu\text{g/ml}$. This

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is consistent with a study by Mahdi *et al.* (2024), which found that the purified L-glutaminase had significant antimicrobial activity against isolates of *P. aeruginosa* ($p < 0.05$) and that it was more effective than the referral standard drug at preventing the formation of biofilms. According to Zhang *et al.* (2005), 28% of the *Lactobacillus* sp. isolates that exhibited enzyme activity also exhibited antibacterial activity. Elborai *et al.* (2023) reported that L-glutaminase shown efficacy against harmful microorganisms. According to Mahdi *et al.* (2019), Amikacin at sub-MIC dramatically reduced the production of *P. aeruginosa* biofilms. Mahdi *et al.* (2021) reported using sub-MICs of gentamicin to minimize the growth of biofilms over plastic sheets.

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فحص وتشخيص وإنتاج إنزيم L-glutaminase بواسطة بكتريا *Lactobacillus reutri* المعزولة من براز البالغين الأصحاء وتأثيره على إصابة الجروح ببكتيريا *P.aerogenosa*

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المستخلص

ان *Lactobacillus reutri* لديها القدرة على ان تستخدم كتأثير دوائي ضد البكتريا المرضية في المستقبل. وتم تحسين التأثير المضاد للميكروبات لـ L-Glutaminase المعزول من *L. reutri* الذي عزز إنتاج الإنزيم في ظل الحد الأدنى من وسائل وطرق الإنتاج. و تكمن أهمية هذه الدراسة في تعزيز إنتاج L-Glutaminase المعزول من *L. reutri* بنشاط عالي باستخدام-L-Glutaminase كمضاد حيوي ومضاد للاغشية الحيوية للبكتيريا. في كروماتوغرافيا التبادل الأيوني، كان النشاط النوعي هو 4.7 وحدة / ملغ من البروتين، مع إنتاجية 65% و 6.6 purification folds. اما في الترشيح الهلامي، كان النشاط النوعي 10 وحدة/ملغ بروتين، مع إنتاجية 60% و 16.6 purification folds ، ووجد أن الوزن الجزيئي لانزيم L-Glutaminase يتراوح بين 34-40 كيلو دالتون تم تقديره بطريقة Sodium Dodecyl Sulfate-Polyacrylamide. وفقا للنتائج، أظهر L-Glutaminase المعزول من *L.reutri* المنقى نشاطاً مضاداً للميكروبات بشكل كبير ضد عزلات *P. aeruginosa* ($P < 0.05$)، وكان نشاط تكوين الغشاء الحيوي للـ L-Glutaminase المنقى أقوى من نشاط المضاد الحيوي للدواء القياسي المرجعي، ($P < 0.05$). Pipracillin tazobactam. قد

يكون هذا اكتشافاً واعداً لعلاجات الأمراض البكتيرية المستقبلية مثل البروبيوتيك الطبيعي.