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JCHR (2023) 13(5), 739-746 | ISSN:2251-6727



# Purification and Characterization of Xylanase Enzyme from Bacillus Substilis Isolated from Forest Soil

### Arifa. P.P, Nirmala Devi. N\*, K. Baskaran, Haseera. N.

Department of Biochemistry, Sree Narayana Guru College, K.G. Chavadi, Coimbatore-641 105.

\*Correspondence Author: Dr. Nirmala Devi, Msc, MPhil, PhD.,

Associate Professor and Head, Department of Biochemistry, Sree Narayana Guru College, Coimbatore-641105, Tamilnadu, India,

(Received: 02 S	September 2023 Revised: 14 October	Accepted: 07 November)		
<b>KEYWORD:</b>	ABSTRACT:			
Bacillus substilis,	The xylanase obtained from Bacillus substilis	isolated from forest soil was purified and		
dialysis,	characterized to assess its future in industrial	application. The purification process was by		
ion-exchange	ammonium sulphate fractionation followed by d	ialysis, ion exchange chromatography and gel		
chromatography,	filtration chromatography. Solid ammonium	sulphate salt was used in concentration of		
SDS PAGE,	80%. The precipitated protein was recovered by	centrifugation and dissolving it in minimum		
Zymogram	quantity of buffer after which the protein were retrieved by dialysis. Xylanase further purified			
	by DEAE cellulose ion exchange column. The	yield and fold purification was $60.47\%$ and		
	2.45% with specific activity of 7.495 U/mg prote	in. Column chromatography using Bio-Gel P-		
	60 matrix indicate high purity yield of 53.46%	and fold purification 6.25% with specific		
	activity of 19.08 U/mg protein. The molecular v	weight was found to be 22Kd ,determined by		
	SDSPAGE .Zymogram analysis of xylanase acti	vity using purified protein resulted in a single		
	band. There was a gradual increase in the er	zyme activity from 2% to 3% of substrate		
	concentration. The enzyme showed stability at a	kaline pH7 and showed an increase in activity		
	at 50°C.The xylanase activity was strongly inl	nibited by Hg2+ and Cu2+ and enhanced in		
	presence of Mn2+.			

### INTRODUCTION

Xylanase are class of enzymes that degrade linear polysaccharide xylan in to xylose. Industrial interest in xylanases is for biofuel production, chemical and pharmaceutical industries, wood pulp bioleaching, paper making, the manufacture of food and beverages, and animal nutrition. The xylanase producing bacterial isolate was collected from soil of forest areas and dense plantation and screened on Birchwood xylan Agar media using Congo red assay If a new efficient xylanaseproducing microorganism is isolated, it is essential to purify and characterize the enzymes to know the action towards substrates of each component of a xylanolytic complex, its regulation and biochemical properties in order to develop more competitive processes. The xylanase producing bacterial isolate Bacillus substilis was collected from soil of forest areas and dense plantation and screened on Birchwood xylan Agar media using Congo red assay

Enzyme purification is a complex and tedious process which involves a combination of methods but this step is far more important because enzymes should be available in their purified form if they are to be used for further studies or lest they should be employed for industrial applications. The choice of a suitable method ultimately depends on the nature of enzyme to be purified. In the present study, efforts were made to purify the xylanase enzyme produced from *Bacillus subtilis*. Crude enzyme extract (1000 ml) was prepared by growing the bacterial isolates independently under their respective optimized conditions. The purification process was initiated by ammonium sulphate fractionation followed by dialysis, ion exchange chromatography and gel filtration chromatography

### MATERIAL AND METHODS PURIFICATION AND CHARACTERIZATION OF XYLANASE

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# Calculation of yield of protein, yield of enzyme activity, fold of purification

Yield of protein and enzyme activity of each fraction obtained during purification is the percentage activity obtained by dividing the total protein content or activity of that fraction with the total protein content or activity of the crude extract as the case may be.

Yield of Protein= (Total protein content of the fraction)/(protein content of the crude extract ) x100

Yield of activity= (Total activity of the fraction)/ (Total activity of the crude extract) x100

Specific activity= (Enzyme activity (U/mL))/ (mg of protein /mL) x100  $\,$ 

Fold of Purification= (Specific activity of the fraction)/ (Specific activity of the crude extract) x100

### **PROTEIN ESTIMATION**

Protein content was estimated according to the method of Lowry *et al.*, 1951 using bovine serum albumin (BSA) as the standard.

### ESTIMATION OF ENZYME ACTIVITY

Xylanase assay was done to determine the activity of the enzyme xylanase. The xylanase enzyme activity was done by measuring the reducing sugar released by the reaction on the birch wood xylan. Thus, the xylanase assay was done according to 3, 5 – dinitro salicylic acid (DNS) method.

The amount of enzyme produced by each isolate in liquid xylan medium was found. 1.0% of the birch wood xylan was dissolved in 50 mM Glycine- NaOH buffer (pH -9.2) and this was used as substrate. 0.5 mL of the buffered substrate (1.0% birch wood xylan and 50mM Glycine NaOH buffer) was reacted with 0.5 mL of crude xylanase enzyme at a temperature of 55°C. The reaction was stopped after 10 minutes by adding 3 mLDNS reagent and then kept in a boiling water bath for 5 minutes. After cooling for a few minutes, the released xylose was quantified at 540 nm against a reagent blank. A reagent blank was made in the same manner except the crude enzyme was not added and 0.5 mL buffer was added. An enzyme blank was also made in which the reagent was added before the addition of enzyme so that only the reducing sugar is estimated. A standard of xylose (reducing sugar) was prepared using stock concentration 1 mg/mL. One unit of xylanase enzyme activity is defined as 1 µmole of xylose liberated per minute per mL of enzyme preparation under standard

assayconditions.

### Standard Curve:

 $\Delta$ A540nm Std = A540nm Std - A540nm Std Blank Prepare a standard curve by plotting the  $\Delta$ A540nm Standard vs the µmoles of Xylose.

Sample Concentration Determination:

 $\Delta$ A540nm Sample = A540nm Test - A540nm Blank Determine the µmoles of xylose using the Standard Curve (µmoles of xylose liberated) x (df)

 $(\mu \text{ rrades of xylose} | berated ))((10) x(0.5))$ U| rrd =

df = Dilution factor. 10 =Time of assay (in minutes) as per Unit Definition

0.5 = Volume (in millilitres) of enzyme used units

### AMMONIUM SULPHATEPRECIPITATION

Enzyme production was carried out using submerged state fermentation under optimized conditions. Extraction of the enzyme was performed by ammonium sulphate precipitation method. Salting out by ammonium sulphate is the best-known method for concentration and purification of the enzyme. Ammonium sulphate precipitation was done according to Englard and Seifter, 1990. Crystal ammonium sulphate (Sigma Aldrich, Bengaluru) required to precipitate xylanase was optimized by its addition, at 80% saturation to the crude extracts.

The addition of ammonium sulphate was done under constant stirring at 4°C for 30 min and then stirring was continued overnight at 4°C. Precipitated protein was collected by centrifugation at 10000 rpm for 15 min at 4°C. The precipitate was suspended in a 25 mM Naphosphate buffer (pH-7.4).

### DIALYSIS

The precipitate obtained after ammonium sulphate precipitation was further dialyzed against (Sigma Aldrich, Bengaluru) 25 mM Na-phosphate buffer (pH-7.4). The buffer was stirred gently using a magnetic stirrer to enhance solute exchange. Dialysis tube (Sigma-Aldrich) was first treated to remove the humectants and protectants like glycerin and sulphur compounds present in it and to make the pores of the tube more clear. The pre-treatment involved washing of the tube in running water for 3-4 hours, dipping in 0.3 % (w/v) sodium sulphide at 80°C for 1 min, further washing with hot water (60°C) for 2 min followed by acid wash in 0.2 %

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(v/v) sulphuric acid. Finally the tube was rinsed with distilled water. Dialysis was conducted overnight and the buffer was changed several times to increase the efficiency of the dialysis. The precipitated protein was dialyzed in the pre-treated dialysis tube for 48 hours at 4°C with several changes of buffer and assayed for xylanase activity, protein content, and specific activity. After dialysis, the sample was collected and stored under-20°C.

### ION EXCHANGECHROMATOGRAPHY

Purification of xylanase from bacteria was carried out by the DEAE-cellulose (Sigma Aldrich, Bangalore) column chromatography. The column was packed with activated DEAE- cellulose equilibrated with 25 mM sodium phosphate buffer (pH 6.0). The height of the column was 20 cm with a 2.5cm diameter. The protein was eluted with the 0.1 to 0.5 M NaCl gradient. The 25 fractions were collected having 3 mL volume of each fraction with the flow rate of 1mL/min. All the steps were carried out at 4°C to prevent proteindenaturation.

### GEL FILTRATION CHROMATOGRAPHY

Purification of xylanase was also carried out by Bio Gel P-100 (Bio Rad, India). The column was packed at the height of 60 cm having a 1 cm diameter and activated using 50mM TrisHCl buffer, pH 7.5. The void volume was calibrated by using blue dextran. The experiment was performed at the flow rate of 1mL/10 min. 25 fractions (1 mL each) were collected and analysed xylanase activity. All the procedures were carried out at4°C.

# MOLECULAR WEIGHT DETERMINATION BY SDSPAGE

Purity of the fractions, showing xylanase activity, was checked by SDS- PAGE by the method of (Laemmli, 1970). The molecular weight of xylanase was determined by comparison with standard molecular marker proteins (Promega Corporation,USA).

### ZYMOGRAM ANALYSIS

Zymography is a technique used to assess the enzymatic activity of proteins either in situ or by separating them with electrophoresis. The enzyme converts the substrate into a product which is detected by different staining methods. The zymogram analysis was performed by loading the concentrated partially purified xylanase fraction on a SDS- 12 % polyacrylamide gel containing 0.1 % xylan. After electrophoresis, the gel was incubated for 60 min at 80 C, soaked in 0.1 % Congo red solution for 30 min at room temperature and washed with 1 M NaCl until excess dye was removed. On gel analysis, the activity band was seen as a clear colourless area of xylan hydrolysis

### CHARACTERIZATION OF PURIFIED XYLANASEENZYME

**Effect of substrate concentration on xylanaseactivity** The substrate specificity of the enzyme on xylanase activity was measured by considering different xylan substrate concentrations in Glycine-NaOH buffer (pH7.0) such as 0.5% - 4% at 37°C.

#### Effect of temperature on xylanaseactivity

The effect of temperature on xylanase activity was studied by incubating the enzyme at a range of  $20^{\circ}$ C to  $60^{\circ}$ C; using 3% (w/v) xylan as substrate in Glycine-NaOH buffer (pH7.0). Optimal temperature was defined as the temperature at which maximum hydrolysis occurred, detected by increase in activity.

### Effect of pH on xylanase activity

The effects of pH on enzyme activity was determined over a pH range of 3.0 to 10.0; using 1 mL of 3% (w/v) xylan as the substrate prepared in citrate- phosphate buffer (pH 3.0-5.0), sodium phosphate buffer (pH6.0-8.0) and glycine- NaOH buffer (pH 9.0- 10.0). The enzyme suspension was incubated for 3 hours in various buffers having pH ranging from 3.0 to 10.0 at 50°C. Optimal pH was defined as the pH at which maximal enzyme activity was obtained in theassay.

#### Effect of metal salts on xylanaseactivity

Effect of various metal ions on enzyme activity was evaluated by incubating the enzyme along with different concentrations of 1mM of various metal salts in the enzyme reaction mixture for 1 hour. The metal salts studied included calcium chloride, mercuric chloride, ferric chloride, copper sulphate, manganese chloride, magnesium sulphate, zinc sulphate, which contributed the metal ions  $Ca^{2+}$ ,  $Hg^{2+}$ ,  $Fe^{2+}$ ,  $Cu^{2+}$ ,  $Mn^{2+}$ ,  $Mg^{2+}$  and  $Zn^{2+}$  respectively.

#### Analysis of Xylanase Activity

Xylanase assay was done to determine the activity of the enzyme xylanase. The xylanase enzyme activity was

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done by measuring the reducing sugar released by the reaction on the birchwoodxylan . Thus, the xylanase assay was done according to 3, 5 – dinitro salicylic acid (DNS) method. The amount of enzyme produced by each isolate in liquid xylan medium was found.

### **RESULTS AND DISCUSSION**

### **Ammonium Sulphate Precipitation**

Precipitation of the enzyme did occur when the crude extract obtained from *Bacillus subtilis* was treated with 80% ammonium sulphate salt concentration



Fig:1. Ammonium Sulphate Precipitation from crude extract

### Dialysis

The crude enzyme obtained after ammonium sulphate precipitation was further purified by dialysis



Fig:2. Dialysis of precipitated protein from ammonium sulphate precipitation

### Ion exchange chromatography

Xylanase was further purified by the DEAE cellulose ion exchange column. The enzyme was eluted from DEAE cellulose column at a NaCl concentration of 0.1 to 0.5 M. The fractions having maximum specific activity were concentrated. The yield and fold purification of Xylanase enzyme was 60.47% and 2.45% respectively with specific activity of 7.495U/mg protein.

### **Gel Filtration**

Dialysed xylanase enzyme obtained from dialysis with Sephadex G-100 matrix was purified by column chromatography using Bio-Gel P-60 matrix where enzyme was eluted out as a single sharp homogenous peak, as revealed from the chromatogram, indicating high purity. Xylanase activity was eluted at about 40-50 fractions using the sodium phosphate buffer (0.05 M, pH 7) as the eluent. The active fraction were pooled and concentrated by ammonium sulphate (80% precipitation) •and dialysed for 24 hrs and considered as pure xylanase for further studies. The yield and fold purification of Xylanase enzyme was 53.46% and 6.25 % respectively with specific activity of 19.08 U/mg protein. The purification reveals the presence of multiple isoforms ofxylanase

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**Table:1.** Showing Total protein, Total Enzyme activity, Specific activity, purification and yield of Xylanase by crude

 extract, Ammonium sulphate precipitation and dialysis, column chromatography and ion exchange chromatography

Xylanase enzyme	<b>Total Protein</b>	Total enzyme	Specific activity(U/mg	Fold of	% yield of
	(mg)	activity (U)	of protein)	purification	enzyme
Crude extract	1132.5	3452	3.049	1.00	100
Ammonium Sulphate	549	2746	5.001	1.63	79.54
Precipitation and dialysis					
Ion exchange	278.5	2087.5	7.495	2.45	60.47
chromatography					
Gel filtration	96.7	1845.5	19.08	6.25	53.46

### Molecular Weight Determination by SDSPAGE

The SDS page was carried out to determine the molecular weight of the purified protein. A clear band was observed in the final purified product. The molecular weight was determined with the help of a protein marker and was found to be approximately 22kD



Fig:3. SDS-PAGE Lane 1: Protein Marker Lane 2: 22kDa purified xylanase protein

### ZymogramAnalysis

Zymogram analysis of xylanase activity using purified protein from *Bacillus subtilis*was carried out. Staining of

the polyacrylamide gel with Congo red resulted in a single band.



Fig:4. Zymography analysis of Xylanase showin clear zone

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### Characterization of enzyme

#### Effect of substrate concentration onactivity

The effect of various substrate concentrations was determined in values ranging from 0.5% - 4% on the purified xylanase activity. Enzyme activity was found to

be greatly affected by substrate concentration. The enzyme exhibited a gradual increase in enzyme activity from 2% to 3% (111.3 to 131.1U/ml) of substrate concentration, respectively and then decreased.



Graph: 5. Effect of Substrate Concentration on Xylanase Activity

#### Effect of temperature on activity

Measurement of the activity of purified extracellular xylanase from *B. subtilis* at different temperatures showed an increase in activity up to 50 °C followed by a

decline. Hence, the optimum temperature for purified *B*. *subtilis* extracellular xylanase was 50 °C.



Graph: 6.Effect of Temperature on Xylanase Activity

### Effect of pH on activity

was observed at this pH.

The optimum pH of purified extracellular xylanase from *B. subtilis*was found to be 7.0 since maximum activity

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Graph:7. Effect of pH on Xylanase Activity

### Effect of metal salts on activity

Xylanase activity was strongly inhibited by Hg2+ which might be due to its interaction with sulfhydryl groups present on the enzyme. Cu2+ ions were also found to be inhibitory in enzyme activity. Some other metal ions like Ca2+, Fe3+, Cu2+, Mg2+and Zn2+ decreased the enzyme activity but to a very less extent. In contrast, xylanase activity was enhanced in the presence of Mn2+ (138.3 U/ml).



Graph: 8. Effect of Metal ions on Xylanase Activity

### CONCLUSION

The xylanase enzyme isolated can be purified by ammonium sulphate followed by dialysis, ion-exchange chromatography. The yield and fold purification of Xylanase enzyme was 60.47% and 2.45% respectively with specific activity of 7.495U/mg protein.Gel filtration chromatography reveals the presence of multiple isoforms of xylanase. The yield and fold purification of Xylanase enzyme was 53.46% and 6.25 % respectively with specific activity of 19.08 U/mg protein. molecular weight was found to be approximately 22kD by SDS PAGE. Enzyme activity was found to be greatly affected by substrate concentration. The enzyme exhibited a gradual increase in enzyme activity from 2% to 3% (111.3 to 131.1U/ml) of substrate concentration, respectively and then decreased. Effect of temperature showed an increase in activity up to 50 °C followed by a decline. The maximum activity was observed at Ph7.metal salt activity showed that Xylanase activity was strongly inhibited by Hg2+ and was enhanced in the presence of Mn2+ (138.3 U/ml). The optimized xylanase activity was 138.7U/ml.

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JCHR (2023) 13(5), 739-746 | ISSN:2251-6727



We thank **Dr. D. Kalpana.** Principal, Sree Narayana Guru College, for the facilities provided is gratefully acknowledged. I also thank Zygene Biotechnologies Pvt Ltd, Cochin for the facilities provided.

### CONFLICT OF INTEREST

Authors declare that they do not have any conflict of interest

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