ISOLATION, IDENTIFICATION AND CHARACTERIZATION OF CELLULASE 3 FROM A BACTERIUM OBTAINED AT A SAW-MILL SITE IN ILE-IFE, NIGERIA

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Submission date: 09-Oct-2017 09:11AM (UTC+0500)

Submission ID: 859536602

File name: Ms_BJI_37221.pdf (755.73K)

Word count: 4345

Character count: 21767

1	Original Research Article
2	ISOLATION, IDENTIFICATION AND CHARACTERIZATION OF CELLULASE
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4	FROM A BACTERIUM OBTAINED AT A SAW-MILL SITE IN ILE-IFE, NIGERIA
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6	ABSTRACT
7	This research characterized cellulase with enviable physicochemical parameters from a
8	bacterium isolated from decaying sawdust heap. Isolated bacteria were screened for
9	cellulolysis using the Congo red plate method. The bacterium with the largest halozone was
10	identified by its 16S rRNA sequence. Optimum growth and cellulase production condition
11	was determined by varying incubation time, pH, temperature, different carbon and nitrogen
12	sources, % substrate concentration and inoculum size. Cellulase was extracted, assayed and
13	partially purified. The kinetic parameters were determined as well as the effect of selected
14	conditions on the activity of the enzyme. Seven isolates showed cellulolytic capabilities.
15	Isolate A8 with 58 mm halozone had 96% sequence identity with Bacillus subtilis FJ532063.
16	Optimum activity of 46.18 U/ml at 28 hours was recorded at pH 7, 35 ± 2 °C. Yields of 18.5
17	and 13.5% resulted from ion exchange and gel filtration chromatography respectively. K_m
18	was found to be 0.0108 ± 0.0032 mg/ml with a V_{max} of $119.3 \pm 7.4 \mu mol/min$. Maximum
19	activity for partially purified cellulase was recorded at pH 9.5 and 55 °C with stability at 50
20	$^{\rm o}{\rm C};$ and pH 9, 35 $^{\rm o}{\rm C}$ with stability at 45 $^{\rm o}{\rm C}$ for crude cellulase. The study revealed that nature
21	is full of cellulolytic bacteria that could be exploited for applications in biotechnology such as
22	hydrolysis of under-utilized lignocellulosic material such as sawdust, to glucose which can
23	further act as feedstock for other value adding products.
24	Keywords: Decayed Sawdust, Bacteria Isolate, Cellulase, Bacillus.
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1. INTRODUCTION

Life thrives on a string of biochemical reactions driven by various enzymes. Enzymes are thus a necessity for the continuous existence of the biological world. Cellulose, which accounts for approximately 1.5×10^{12} tons of biomass produced through photosynthesis annually, is the most abundant organic compound on earth (Guo et al., 2008). This abundance has made cellulase enzyme one of the most sought after in the commercial market as it degrades cellulose. Nature is rich in microbial groups with cellulolytic abilities such as fungi, actinomycetes and bacteria. Higher organisms such as insects, arthropods and plants have been found with various degrees of cellulolytic capability (Fischer et al., 2013; Duan and Feng, 2010; Sun and Scharf, 2010). For this study, bacteria have been selected for their profuse growth and shorter generation time when compared with their bio-counterparts. A significant amount of diversity exists among cellulolytic bacteria. Bacteria cells are sources of cellulase irrespective of the gram reaction, oxygen requirements or other basis of classification. Various Gram negative, Gram positive and Gram-variable bacteria produce cellulase (Yan and Wu, 2013; Huang et al., 2012). Cellulolytic bacteria could also be aerobic, facultatively anaerobic or anaerobic (Bayer et al., 2007). Cellulolytic bacteria have been isolated from a wide diversity of environments; extreme or favourable. Acidothermus, Bacillus, Clostridium, Pseudomonas, Rhodothermus, Microbacterium, Rhizobioum and Escherichia are genera that have been exploited for cellulase production (Huang et al., 2012; López-Contreras et al., 2004; Heck et al., 2002). This study aimed to find a cellulolytic bacterium capable of producing active cellulase in substantial amounts and with enviable physicochemical parameters from a saw mill.

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2. MATERIALS AND METHOD

54 2.1 Sample collection and preparat	tion
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- 55 Unweighed quantity of sawdust was collected into a sterile bottle from a decaying sawdust
- heap at the saw mill located at Modakeke, Ile-Ife at a depth of about 2 metres. One gram (1 g)
- 57 of decaying sawdust was accurately weighed out and dispensed into 10 ml sterile distilled
- 58 water in a test tube. It was mixed well to ensure even dispersal of the microbial flora in the
- 59 sample. This made the stock preparation. Aliquots of 1 ml was aseptically pipetted from the
- stock and transferred into the next tube of 9 ml sterile distilled water and mixed properly.
- This made the 10^{-1} dilution. The procedure was repeated until the sixth tube (10^{-6} dilution).

62 2.2 Bacteria Isolation

- Aliquots of 1 ml of 10⁻⁴, 10⁻⁵, 10⁻⁶ dilution were plated out in duplicates using pour plate
- 64 technique. Pure cultures were subsequently obtained and stored for further use.

65 2.3 Screening for Cellulolytic Ability

- 66 Carboxymethylcellulose agar (CMCA) plates incubated with a single streak of pure isolate
- 67 were flooded with 0.1% Congo red solution after 48 h and de-stained with 1 M NaCl
- 68 solution. A clear halozone around the line of streak depicted cellulose hydrolysis. The
- 69 diameter of the halozone was measured and the isolates with considerable large halozones
- 70 were picked for further studies.

71 2.4 Bacterial Identification

- 72 Pure cultures of cellulolytic bacteria were identified by their reactions to biochemical tests
- 73 and the strain with maximum cellulase activity was further subjected to molecular
- 74 identification by an analysis of the 16S rRNA sequence.

76 2.5 Submerged Fermentation Process

- 77 The cellulolytic bacterial cultures were grown over a period of 48 h in 0.1 M Phosphate
- buffer, pH 7.0 containing bacteriological peptone (2% w/v), K₂HPO₄, (0.3% w/v),
- 79 MgSO₄.7H₂O (0.1% w/v), NaCl (0.075% w/v) and high viscosity carboxymethylcellulose
- 80 (0.2% w/v) as depicted by Kotchoni and Shonukan, 2002. This was done with agitation at
- 81 150 rpm in a water bath shaker. The medium was continually assayed for cellulase every 2 h.
- Thus, the best cellulolytic bacterium alongside its growth pattern was determined.

83 2.6 Cellulase Extraction

- The growth medium, after optimal incubation, was centrifuged at 12, 000 rpm for 20 minutes
- and at a cold temperature of 4°C. The supernatant was used as the crude enzyme.

86 2.7 Cellulase Assay

- 87 Cellulase activity was measured by the presence of reducing sugars released by the
- 88 hydrolysis action of the enzyme on its substrate using Somogyi-Nelson method (Somogyi,
- 89 1952; Nelson, 1944). The reducing sugars were determined by incubating 0.1 ml of 0.2% w/v
- 90 CMC, stabilized by 0.80 ml 0.1 M phosphate buffer, pH 7.0 with 0.05 ml of crude enzyme
- 91 and inactivated crude enzyme (boiled at 100°C for 15 minutes) at 37°C for 20 mins. The
- 92 reaction was terminated by the addition of 1 ml alkaline copper tartrate solution and
- 93 subsequent boiling for 20 minutes. One millilitre (1 ml) of arsenomolybdate solution was
- 94 added after cooling for colour stabilization. Absorbance was read at 540 nm against a reagent
- 95 blank by a spectrophotometer and the amount of reducing sugars was interpolated from the
- 96 glucose standard curve.

97 2.8 Optimization of Cellulase Production Conditions

- 98 The pH, temperature, carbon source, nitrogen source, percentage substrate concentration, and
- 99 inoculum size of the basal medium was varied to observe the effect on enzyme production.

100	pH was varied from 4-10; temperature from 30-60°C; carbon sources (glucose, sucrose,
101	lactose, maltose, galactose and mannitol); nitrogen sources (tryptone, yeast extract, malt
102	extract and urea for organic nitrogen; NH ₄ Cl, (NH ₄) ₂ SO ₄ , NH ₄ NO ₃ , NaNO ₃ and NH ₄ H ₂ PO ₄
103	for inorganic nitrogen); percentage substrate concentration was varied from 0.2-1.0% and
104	inoculum size was varied from 1-5%. In each case, all other conditions were held constant.
105	2.9 Cellulase Purification
106	Cell Free Supernatant (CFS) was partially purified by precipitation with 80% ammonium
107	sulphate and acetone and then, dialysis. CFS was also concentrated by lyophilization. Further
108	purification was done by Ion exchange chromatograpy on diethylaminoethyl (DEAE)-
109	Sephacel and Gel filtration chromatography on Sephacryl S-200.
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111	2.10 Determination of Kinetic Properties
112	Kinetic parameters (K_m and V_{max}) were determined for the partially purified cellulase by
113	incubating aliquots of the enzyme with CMC to make a final substrate concentration in the
114	range 0.01-0.1 mg/ml and estimating the sugars released. Conditions for cellulase activity
115	were optimized.
116	2.11 Effect of Temperature, pH and Heat Stability on crude and Partially Purified
117	Cellulase
118	Aliquots of the enzyme was incubated with substrate and reducing sugars
119	estimated as depicted in 2.7 at varying conditions of temperature (30-60 °C),
120	pH 4 - 10 and 35-70 °C for heat stability.
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3. RESULTS AND DISCUSSION

124 3.1 Bacteria Isolation and Characterization

125 As shown in Table 1, Bacillus cereus, B. subtilis, B. brevis, B. circulans, Serratia

marcescens and B. megaterium were the cellulolytic bacteria isolated as A3&A21,

A8, A11, A13, A15 and A22 respectively, as compared with the Bergey's Manual

128 of Determinative Bacteriology.

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Table 1: Morphological and Biochemical Characteristics of Cellulolytic Isolates

Isolate code	A3	A8	A11	A13	A15	A21	A22
Halozone Diameter (mm)	47	58	26	21	32	25	39
Morphological Characteristics							
Gram reaction	+	+	+	+	-	+	+
Shape	Rods						
Spore Staining	+	+	+	+	ND	+	+
Motility	+	+	+	+	+	+	+
Biochemical Characteristics							
Catalase	+	+	-	-	+	+	+
Citrate	+	+	-	-	+	+	+
Starch Hydrolysis	+	+	+	+	ND	+	+
Methyl Red	+	-	-	-	-	+	+
Voges Proskauer	+	+	-	-	+	+	-
Nitrate Reduction	+	+	ND	ND	+	+	+
Growth in 6.5% NaCl	ND	+	-	+	ND	ND	+
Oxidase	+	ND	-	ND	-	+	ND
Indole	-	-	-	-	-	-	-
Sulphide	-	ND	-	-	-	-	ND
Urease	-	ND	-	ND	-	-	-
Sugar Utilization							
Glucose	+	-	+	-	+	+	+
Lactose	+	-	-	+	-	+	+
Mannitol	-	+	-	+	+	-	+
L-arabinose	-	ND	-	+	-	-	-

131 Keys: + = positive reaction, - = negative reaction, and ND = Not Determined

132 Approximately, 85.7% of the isolates were identified as Bacillus species. This shows the

predominance of *Bacillus* species as organisms of interest in cellulase production.

3.1 Screening for Cellulolysis

Diameter of halozones recorded vary among the isolates. A8 had the largest diameter of 58 mm (Fig 3.1).



Fig. 3.1: Halozone of Isolate A8

3.2 Molecular Identification

Isolate A8 was further found to have a 96% similarity with the rRNA sequence of B. subtilis with the accession number FJ532063 of the GenBank, hence the isolate was confirmed as *Bacillus subtilis* A8.

3.3 Optimum Conditions for Cellulase Production from B. subtilis A8

As depicted in Fig 3.2, the growth pattern of *B. subtilis* A8 revealed a lag phase of about 6 h; logarithmic phase of about 28 h; stationary phase of about 6 h. The peak of cellulase activity was however at 36 h, falling in the stationary phase $(34^{th} - 40^{th})$ hour). This confirms enzymes as secondary metabolites. This however was in contrast with the findings of Shabeb *et al.* (2010) and Mukesh Kumar *et al.* (2012) where maximum cellulase productivity from *B. subtilis* was recorded after 24 h and 72 h respectively. Maximum activity of cellulase at 36^{th}

hour of incubation is of a better advantage. This is because equipment and facilities are tied down in use for shorter periods, lower cost is incurred and less energy consumed.

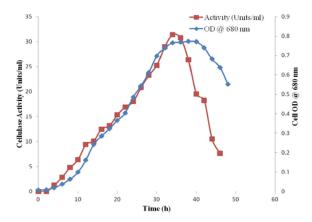


Fig. 3.2: Growth and cellulase activity of Bacillus subtilis A8 at 37°C, pH 7

Just as reported by Jayadev, 2014, *B. subtilis* A8 showed the highest cellulase activity at pH 7 (Fig. 3.3). The trend observed however, showed a preference for alkaline over acidic medium. On the contrary, Vijayaraghavan and Vincent (2012) reported the preference of a *Bacillus* sp. for a slightly acidic medium, with optimum pH at 6.5 while there was very low activity at pH 8.0.

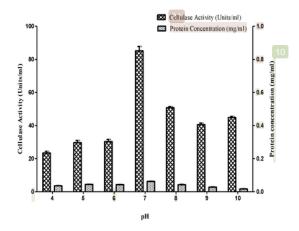


Fig. 3.3: Effect of pH on cellulase production by Bacillus subtilis A8 at 37°C

An optimum temperature of $35 \pm 2^{\circ}$ C was recorded in this present study (Fig. 3.4). This same optimum temperature was reported by Amritkar *et al.* (2004). At this relatively low temperature, not much heat is generated hence, there is little or no need for cooling systems in industries, and less energy is consumed. Shabeb *et al.* (2010) however reported maximum cellulase activity by *B. subtilis* at a higher temperature of 45° C.

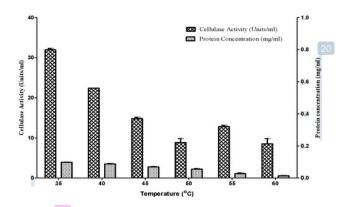


Fig. 3.4: Effect of temperature on cellulase production by Bacillus subtilis A8 at pH 7

The preferred choice of carbon source for *B. subtilis* A8 as shown in Fig. 3.5 was CMC and not any of the simple sugars used. This is of benefit in the industrial production of cellulose, as many low-cost carbon sources employed contain carbon in complex forms.

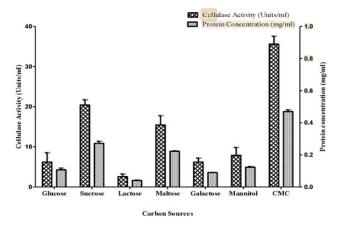


Fig 3.5: Effect of various carbon sources on cellulase production by Bacillus subtilis A8 at 37°C, pH 7

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Of the nitrogen sources tested, the organic nitrogen sources supported growth and cellulase production better than the inorganic nitrogen sources (Figs. 3.6). This is of immense benefit as organic nitrogen sources abound more in nature.

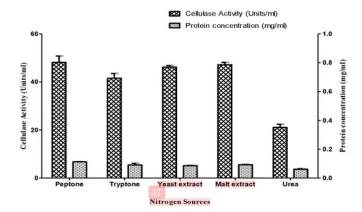
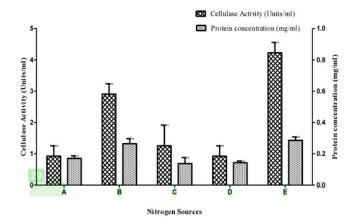


Fig. 3.6a: Effect of various organic nitrogen sources on cellulase production by Bacillus subtilis A8 at 37°C, pH 7



180 Keys: A - NH₄Cl; B - NH₄NO₃; C - (NH₄)₂SO₄; D - NH₄H₂PO₄; E - NaNO₃

Fig. 3.6b: Effect of various inorganic nitrogen sources on cellulase production by Bacillus subtilis A8 at 37°C, pH 7

Findings by Gautam *et al.* (2010) showed a maximum yield of cellulase at 1% concentration, a higher figure than that obtained in this study (Fig. 3.7). A lower substrate concentration is

of good economic value to industries and also individual researchers, as it reduces the cost of production.

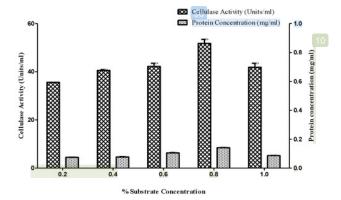


Fig. 3.7: Effect of percentage substrate concentration on cellulase production by Bacillus subtilis A8 at 37°C, pH 7

The highest cellulase activity was obtained from the use of 4% inoculum size. This contrasts with the 8 and 10% recorded by Omojasola and Jilani (2009) and Fadel (2000) respectively. A lower inoculum size is better for less competition for resources by the organisms, thereby increasing the production of metabolites.

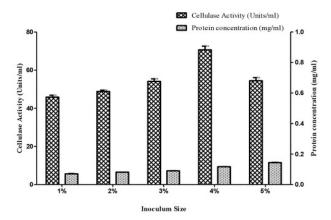


Fig. 3.8: Effect of percentage inoculum size on cellulase production by Bacillus subtilis A8 at 37°C, pH 7

Acetone and ammonium sulphate precipitation were partial purification methods employed, both of which resulted in a considerable loss of activity, hence, the decision to lyophilize (Table 2). Concentration of the CFS by lyophilization considerably shortened the time involved in the purification process as there was no further need for dialysis.

Comparison of Partial Purification Methods for Cellulase from Bacillus subtilis A8

PROCEDURE	VOLUME (ml)	ACTIVITY (Units/ml)	TOTAL ACTIVITY (Units)	PROTEIN (mg/ml)	TOTAL PRROTEIN (mg)	SPECIFIC ACTIVITY (Units/mg)	YIELD (%)	PURIFICATION FOLD
Crude	30	46.18	1385.40	7.24	217.20	6.38	100	
80% Ammonium Sulphate Precipitation	9	61.20	550.80	9.63	86.67	6.36	39.76	1.00
Acetone Precipitation	4	70.45	281.80	7.29	29.16	9.66	20.34	1.51
Pre- Lyophilized	40	46.18	1847.20	7.24	289.60	6.38	100	
Lyophilized	5	334.46	1672.30	11.08	55.40	30.19	90.53	4.73

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Purification on DEAE - Sephacel resulted in two broad peaks as shown in Fig 3.9, with the second peak having a higher cellulose activity than the first. This probably represents different components of the cellulase complex.

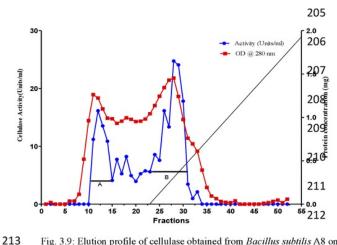


Fig. 3.9: Elution profile of cellulase obtained from Bacillus subtilis A8 on DEAE-Sephacel ion exchange column

214 Further purification by gel filtration on Sephacryl S-200 resulted in a single peak (Fig. 3.10).

A yield of 87.8% recorded from the lyophilized cellulase as shown in Table 3. This implies a

good suitability for cellulose hydrolysis. A lower yield was however recorded from cellulase

partially purified by the chromatographic methods employed. 217

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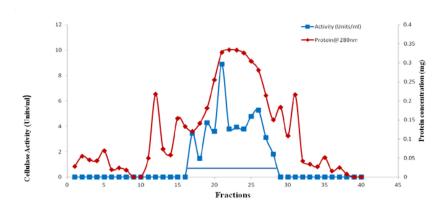


Fig. 3.10: Elution profile of cellulase obtained from Bacillus subtilis A8 on Sephacryl S-200 gel filtration column

	Table 3: Summary of the Purification Protocol of Cellulase Obtained from Bacillus subtilis A8							
PROCEDURE	VOLUME (ml)	ACTIVITY (Units/ml)	TOTAL ACTIVITY (Units)	PROTEIN (mg/ml)	TOTAL PRROTEIN (mg)	SPECIFIC ACTIVITY (Units/mg)	YIELD (%)	PURIFICATION FOLD
Crude	50	50.13	2506.28	7.15	357.54	7.00	100	-
Lyophilized	10	220.05	2200.50	12.13	121.30	18.14	\$7.8	2.59
0 M pooled ion exchange fractions	14.1	8.32	117.31	1.38	19.46	6.03	4.7	0.86
0.5 M pooled ion exchange fractions	8.6	19.47	167.44	2.15	18.49	9.06	6.7	1.29
Lyophilized pooled ion exchange fractions	5	92.76	463.78	6.57	32.85	14.19	18.5	2.03
Gel filtration chromatography	30	11.29	338.70	4.73	141.9	2.39	13.5	0.34

As calculated from Fig. 3.12, the K_m of partially purified cellulase was found to be 0.0108 \pm 0.0032 mg/ml with a V_{max} of 119.3 \pm 7.4 μ mol/min. The low K_m showed high affinity of cellulase from B. subtilis A8 for the substrate (CMC) whereas the high V_{max} is an indication of the rapidness of its hydrolytic capability of the produced cellulase from B. subtilis A8. Linton and Greenaway (2004) reported a much lower V_{max} of 0.01 μ mol/min and 0.03 μ mol/min for total cellulase obtained from the foregut of Gecarcoidea natalis and Discoplax hirtipes respectively.

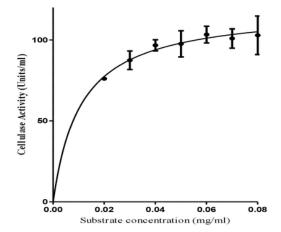
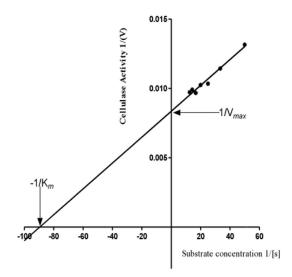


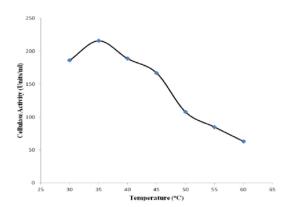
Fig. 3.11: Michealis-Menten plot of partially purified cellulase from Bacillus subtilis A8



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Fig. 3.12: Lineweaver-Burk plot of partially purified cellulase from Bacillus subtilis A8

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Fig. 3.13: Effect of temperature on the activity of crude cellulase obtained from Bacillus subtilis A8 at pH 7

7.0
6.0
(missing) 4.0
5.0
1.0
25 30 35 40 45 50 55 60 65
Temperature (°C)

Fig. 3.14: Effect of temperature on the activity of partially purified cellulase obtained from Bacillus subtilis A8 at pH 7

Crude cellulase had the highest activity at pH 9 (Fig. 3.15) while purified cellulase was more active at pH 9.5 (Fig. 3.16). This is similar to findings by Aygan *et al.* (2011) where optimum for cellulase was pH 10.0. It disagrees with the findings of Yin *et al.* (2010) where maximum cellulase activity from *B. subtilis* YJ1 was recorded at pH 6.0 and that of Linton and Greenaway (2004) which showed an optimum pH of 5.5. Optimum activities at neutral pH values of 7.0 and 7.5 as in the cases of cellulase extracted from *B. coagulans* Co4, *B. amyloliquefaciens* and *Sinorhizobium fredii* have been reported by Adeleke *et al.* (2012).

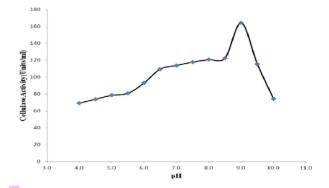


Fig. 3.15: Effect of pH on the activity of crude cellulase obtained from Bacillus subtilis A8 at 35°C

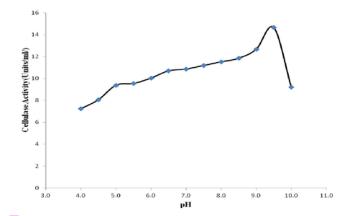


Fig. 3.16: Effect of pH on the activity of partially purified cellulase obtained from Bacillus subtilis A8 at 55°C

Crude cellulase from *B. subtilis* A8 showed high activity and stability at 45°C as depicted in Fig. 3.17. From 55-70°C, there was no significant difference in the level of activity. The relatively low temperature at which the crude cellulase is stable might be as a result of the impurities present.

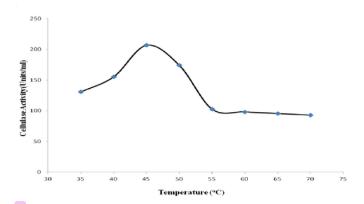


Fig. 3.17: Effect of temperature on the stability of crude cellulase obtained from Bacillus subtilis A8 at pH 9

The enzyme was stable at 50-60°C for at least 60 minutes, retaining 89.67% of its initial activity at optimum temperature (Fig. 3.18). Stability is a necessary characteristic of a good industrial enzyme and cellulase produced from *B. subtilis* A8 showed stability at high temperature. Similar results were found with *Bacillus* sp. CH43 and HR68 (Mawadza *et al.*, 2000).

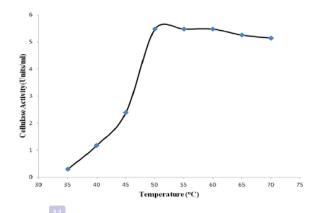


Fig. 3.18: Effect of temperature on the stability of partially purified cellulase obtained from Bacillus subtilis A8 at pH 9.5

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CONCLUSION 265 Bacillus subtilis A8 is a bacterium capable of synthesizing cellulase enzyme with a good 266 hydrolysing capability under mild physicochemical conditions. Also, cellulase from B. 267 subtilis A8 is thermostable with high activity and could therefore be of immense benefit to 268 industries that rely on the use of cellulase. Also, a different resin aside those employed in this 269 study is recommended for better yield. 270 271 REFERENCES 272 Adeleke, E. O., Omafuvbe, B. O., Adewale, I. O. and Bakare, M. K. (2012). Purification and 273 characterization of a cellulase obtained from cocoa (Theobroma cacao) pod-274 degrading Bacillus coagulans Co4. Turkish Journal of Biochemistry, 37(2):222-230. 275 276 Amritkar, N., Kamat, M. and Lali, A. (2004). Expanded bed affinity purification of bacterial 277 α-amylase and cellulase on composite substrate and analogue-cellulose matrices. Process Biochemistry, 39(5):565-570. 278 Aygan, A., Karcioglu, L. and Arikan, B. (2011). Alkaline thermostable and halophilic from 279 280 Bacillus licheniformis C108. African Journal of Biotechnology, 10(5): 789-796. Bayer, E. A., Lamed, R. and Himmel, M. E. (2007). The potential of cellulases and 281 cellulosomes for cellulosic waste management. Current Opinion in Biotechnology, 282 18(3):237-245. 283 284 Duan, C. J., Feng, J. X. (2010). Mining metagenomes for novel cellulase genes. Biotechnology Letters, 32(12):1765-1775. 285 Fadel, M. (2000). Production physiology of cellulases and ß-glucosidase enzymes of 286 287 Aspergillus niger grown under solid state fermentation conditions. Online Journal of Biological Science, 1(5):401-411. 288

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