Purification and Characterization of - Amylase from Bacillus subtilis Isolated from Cassava Processing Sites

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> This study was designed to purify and characterize of -amylase from pure strain of **Bacilla Struck strain strain strain** substitle and an analyze substitled and an analyze substitled and an analyze substitled and an analyz amylase was purified by ammonium sulphate precipitation, then loaded on DEAE Sephadex A-50 ion exchange chromatography and gel filtration. The effect of pH, temperature and metal ions were investigated on the purified enzyme. The single protein band on SDS-PAGE suggested that the enzyme was homogenous. Two different activity peaks were observed in ion exchange chromatography designated pool A and pool B with the 8% and 4% yield, 15.93 and 6.44 purification fold and specific activity 2.55 μmol/min/mg and 1.03 μmol/min/mg respectively. The two fractions revealed the same optimum pH 7.0 for the -amylase activity while the enzyme was relatively stable at pH 4.0 and 7.0 between 20 to 40 minutes and 60 to 80 minutes for pool A and pH 8.0 between 40 and 100 minutes for pool B. At 40°C, optimum temperature was reached, and amylase activity was maintained at 75% and 70% temperature stability between 60 to 80 minutes for pool A and B, less than 20%, the residual activity at 60°C and 70°C was recorded. The incubation of \sim -amylase with Na⁺ and Zn²⁺ ions enhanced/activate the enzyme activity correspondingly, Al^{3+} and K⁺ ions exhibited varied degree of inhibition while Ca²⁺ and Hg²⁺ ions caused total inhibition on -amylase activity. The ability of purified -amylase from **Bacilla is under wide range of** temperatures and pH suggests its applications in industries and bioremediation of effluent discharge on food processing sites.

Bacillus subtilis Cassava processing site; -amylase; Durif cation/Characterization

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Globally, indiscriminate dumping or disposal of wastes to the environment a ects both macro-and-microorganisms causing ecological imbalance and displacement from their niche. e survival of microorganisms depended on several factors such as biotic, and abiotics. Soil stands as major reservoir of di erent microorganisms. Biodegradation of wastes on/in soil by the action of microorganisms could prompt a safer environment [1]. Microorganisms isolated from soil stand as potential source of industrial enzymes due to their abundance in nature. However, utilization of these enzymes such as amylases, cellulases, linamarases, mannanase and pectinase etc in our industries are limited due to the poor exploration. Amylases (EC 3.2.1) is a hydrolytic enzyme that catalyze the hydrolysis of amylose and DK-600), and the reaction was terminated by adding 1 ml of 35dintrosalicylic acid (DNSA) reagent tube. e tubes were incubated for another 5 minutes in a boiling water bath for colour development and cooled rapidly. e activity of the reaction mixture was measured against blank at 540 nm e amount of liberated reducing sugars (glucose equivalents) were estimated by the dinitrosalicylic acid (DNS) method of Miller. One unit of -amylase was defined as amount of enzyme producing 1 micromole of glucose equivalent per minute under the experimental conditions.

Protein determination: e protein estimation of the crude extract was determined by Bradford method and Bovine Serum Albumin (BSA) was used as standard Bradford.

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Durification of -amylase from Bacillus subtilis e purification of crude extract (enzyme supernatant) obtained aіer cold centrifugation was done in three stages. In stage one, 400 ml of the supernatant of crude enzyme was precipitated by adding 240 g ammonium sulphate to obtained 60% ammonium sulphate concentration. e concentrates were then cold centrifuged at 6000 rpm for 30 minutes, the precipitate obtained was washed with $100 \,\mathrm{ml}$ phosphate bu er (pH 6.0) and then dialyzed in the same bu ers" In stage two, the partially purified amylase obtained was purified on ion exchange column chromatography containing DEAE (diethylamineethyl) Sephadex A-50 $(20 \times 25 \text{ cm}$, Pharmacia). e eluates obtained from stage two were washed with 300 ml ion-free water, followed by 200 ml 0.01 M Tris-HCl bu er (pH 8.0). e gel was eluted with NaCl. e absorbance of each of the fraction at time interval was measured. In stage three, 2.5 ml of partially purified -amylase was loaded onto a column chromatography (2.5 cm in diameter and 30.0 cm high) using Sephadex G-150 (Pharmacia). Phosphate elution bu er at 50 mM pH 60 was applied with the flow rate 20 ml/hour. A fraction of 50 ml was collected at interval of 30 minutes and the highest

B are shown (Figures 5 and 6). e –amylase from pool A and pool B increased progressively up pH 4 with specific activity 39.17 μmol/min/mg e optimum activity was attained at pH 7.0 . relative activities of pool A and pool B at pH 6.0 was reduced by 34% and 47% and pH 9.0 was reduced by 10% and 48% respectively when compared with the value obtained at $pH 7.0$ e stability of the enzyme was checked at die rent pH ranged from 5.0 to 8.0 at room temperature for two hours. e – amylase activity obtained from pool A was relatively stable between 20 to 40 and 60 to 80 minutes of incubation at pH 40 and 7.0 while -amylase activity obtained from pool B was stable between 40 to 100 minutes of incubation at pH 8.0 respectively. pH of the growth medium plays an important role by inducing morphological change in the nature of microorganism and their enzyme secretion. e decrease in pH beyond optimum might be due to the concomitant alteration in the conformation of the enzyme protein caused by changes in pH of its environment resulting in changes in the three dimensional structure of the enzyme active sites and substrates binding speed to produce maximum product. pH 6.8-7.2 optimum for purified amylase from Bacillus subtilis has been reported [20]. Roy and Rowshanul [16] reported pH 6.0 optimum for purif ed xylanase from *Bacillus cereus*. It was reported that the stability of most plant enzymes decreased significantlmat pH values below 4.0 and above 7.5, whereas, the majority of the corresponding microbial enzymes are rather stable at pH 7.0 [21]. Mukesh et al. [8] reported $$ amylases stability over a wide range of $pH 4$ to 11. e optimum pH 7.0 and above for $-\text{amylase}$ activity from B. subtilis, B. licheniformis $B.$ amyloliquefaciens and thermophilic Anoxybacillus f avithermus has been reported [16,22-26] pH 8.0 optimum has been reported for amylase activity from ermus sp., Bacillus KSM-K38 and Bacillus spp $[27,28]$ Wide range (pH 3.5 to 1.2) optimum pH for $-$ amylases has been reported [7,29-32]. e result obtained from this study agrees with the findings of Agülo˘glu Fincan and Bukhari and Rehman [32,33] who reported pH 7.0 optimum for purified -amylase from Bacillus subtilis from local environment.

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Figure 5: E ect of pH on the stability of purified -amylase for pool A.

E ect of temperature on -amylase activity and stability: Figure 7 shows the e ect of temperature on purified -amylase activity while the temperature stability of the purified -amylase from pool A and pool B are shown in Figures 8 and 9. He -amylase activity increased with increase progressively with incubation temperature until an optimum percentage relative activity was reached at 40°C and 60°C for pool A and pool B respectively (Figure 7).

Figure 7: E ect of temperature on the activity of purified -amylase.

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