

Analysis of Enzymes Activities on Domestic Waste Dump Sites

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Abstract

Introduction: Effects of physico-chemical parameters on microbial dehydrogenases from domestic waste dumpsites were studied.

Methodology: The microorganisms (*S. aureus*, D2 and EDTA, ethanol and butanol) on the microbial dehydrogenase were determined.

Results: V_{max} and K_m of the microorganisms were 10.58 mg Formazan/mg cell dry wt/h, respectively. Calcium ion, Mg

($p < 0.05$) dehydrogenase activities in all the microorganisms studied while Zn^{2+} , Fe^{2+} and EDTA decreased the activities.

Conclusion: V_{max} and K_m of the microorganisms were 10.58 mg Formazan/mg cell dry wt/h, respectively. Calcium ion, Mg to enhance microbial growth; which is essential for the degradation of domestic waste thereby promoting cleaner environment.

Keywords: Dehydrogenase activities; Enzymes; Domestic waste

Introduction

The quality of life on the Earth is linked inextricably to the overall quality of the environment. Wastes were traditionally disposed of in landfills in the past. This traditional mode of waste disposal was publically unacceptable due to the increasing conversion of scarce agricultural lands to dump

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and suspended in the same buffer containing 2 mM EDTA and 1 mM Dithiothreitol (DTT). Cells were ruptured using osmotic shock by Nossal and Heppel [12]. The cells were suspended in 20% sucrose buffer then separated by centrifugation at 4000 rpm for 10 min. The resulting paste was dispersed in acetone at 4°C. Cellular debris and unbroken cells were removed by centrifugation at 4000 rpm for 45 min at 4°C. The supernatant obtained constituted the crude microbial extracts (soluble enzyme fraction) for each microbe.

Purification of enzyme

The enzyme was partially purified from the crude microbial extracts in four steps: ammonium sulfate precipitation, dialysis, sephadex G200 gel filtration chromatography and DEAE-cellulose column chromatography. All the steps were performed at 4°C.

Ammonium sulphate precipitation

The protein sample was allowed to thaw to determine total volume, and was centrifuged at 3000 rpm for 30 min. This was transferred into a beaker containing a stir bar and was placed on a magnetic stirrer. While the sample was being stirred, solid ammonium sulfate crystals were added to bring the final concentration to 60% saturation. (The volume of ammonium sulphate used was equal to the volume of the

Ion-exchange chromatography

The enzyme was further purified using DEAE-cellulose column chromatography. The method used was as described by Yannis [14]. DEAE-cellulose was suspended in 8 vol. of Tris-buffer containing 50 mM of NaCl and kept overnight for equilibration. The column was carefully packed equilibrated with 8 vol. of same buffer containing 0.25 mM of NaCl. Then 3 ml of the partially purified enzyme was diluted to 15 ml and loaded into the column and was washed with appropriate 100 ml of the equilibration buffer. The protein was eluted with 0.25M buffer pH 7.2 and NaCl gradient 0.1-1 M, was passed through the exchanger

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