

## Bloodstream Form of *Trypanosoma evansi* Contains $\beta$ -Galactosidase

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**Abstract:**  $\beta$ -Galactosidase (EC 3.2.1.23) was isolated and characterized from the bloodstream form of *Trypanosoma evansi*. The galactose hydrolase was purified 6.98 fold with a yield of 33.33% by anion exchange chromatography using DEAE- cellulose and size exclusion procedure on Sephadex G- 75 column. The *T. evansi*  $\beta$ -Galactosidase had broad pH and temperature ranges with an optimum of 6.0 and 37°C respectively. Initial velocity studies for the determination of kinetic parameters using 4-methylumbelliferyl- $\beta$ -D-galactoside (MU-Gal) as substrate revealed a  $K_M$  and  $V_{max}$  of 0.08mM and 0.07  $\mu$ mol/min respectively. The enzyme activity was enhanced by cations:  $Fe^{2+}$ ,  $Co^{2+}$  and  $Mg^{2+}$  while  $Ca^{2+}$  was found to inhibit it.

**Key words:**  $\beta$ -Galactosidase • Enzyme • Galactosides • *T. evansi*

### INTRODUCTION

*Trypanosoma evansi*, the causative agent of *surra*, is a hemoflagellate protozoan parasites that infects domestic animals causing animal trypanosomiasis. It has the broadest geographical and host range among the pathogenic animal trypanosomes. The mechanism of pathogenesis is in part due to the action of sialidase, which cleaves erythrocytes surface sialic acids and expose  $\beta$ -Galactosyl residues. These residues are then recognized by  $\beta$ -D- galactose specific lectins on macrophages leading to erythrophagocytosis and subsequently anemia [1-3].

$\beta$ -Galactosidase is an enzyme that catalyses the hydrolysis of  $\beta$ -Galactosides from lectins, glycoproteins and gangliosides. The enzyme has been purified and characterized in *Tritrichomonas foetus* [4] where it is required for mucin breakdown during invasion process [5]. So far, among all the species of trypanosomes,  $\beta$ -Galactosidase has only been described in *T. congolense* [6, 7] and *T. cruzi* [8, 9] but has never been characterized from any of the African trypanosomes. In this paper, we report for the first time the purification and some biochemical characterization of  $\beta$  galactosidase from *T. evansi*.  $\beta$ -Galactosidase from *T. evansi* may play some significant role in the infection biology of the parasite which could be exploited as a target for deciphering a novel and more potent chemotherapeutic agent than the contemporary antitrypanosomal drugs which have been found resistant by the trypanosomes. Moreso, understanding the key agents involved in the

pathogenesis and pathophysiology of the *T. evansi* infection will improve the current strategies in place for the development of vaccine against trypanosomiasis.

### MATERIALS AND METHODS

Five experimentally *T. evansi* (IL-1392) infected rats were obtained from Department Of Parasitology and Entomology, Faculty Of Veterinary Medicine, Ahmadu Bello University, Zaria- Nigeria. All other reagents were purchased from Sigma Chemical Company, USA.

#### Treatment of Experimental Animals

**Isolation of  $\beta$ -Galactosidase from Bloodstream Form of *T. evansi*:** Bloodstream *T. evansi* parasites were isolated from the rats at a cell density of  $10^8$  trypanosomes/ml of blood. The trypanosomes were separated from the red blood cells (RBC) using the Lanham and Godfrey method [10], centrifuged at 10000xg for 10minutes and washed in phosphate buffered saline (PBS), pH7.2. Freezing and thawing with gentle agitation and treatment with 2% Triton X-100 solubilized the parasites and released the enzyme.

**Purification of  $\beta$  Galactosidase from Bloodstream Form of *T. evansi*:** The purification entailed first a micropurification procedure involving the loading of two mls of the crude enzyme onto a pre-equilibrated DEAE cellulose column (2.5cm X 30cm), pH 7.2 and eluted with a linear gradient of NaCl (0.01-0.1M). Twelve, 5ml fractions were collected and assayed for  $\beta$ -Galactosidase and total

Table 1: Summary of Purification protocols of  $\beta$  galactosidase from *T. evansi*

Purification Step	Protein (mg)	Total activity ( $\mu\text{mol/min}$ ) $\times 10^{-3}$	Specific activity ( $\mu\text{mol/min/mg}$ ) $\times 10^{-3}$	Purification fold	Yield (%)
Crude	0.90	84	93.33	1	100
Ion-exchange chromatography (DEAE-cellulose)	0.12	30	250	2.70	35.70
Gel filtration on Sephadex G 75	0.043	28	651.20	6.98	33.33

proteins. The  $\beta$ -Galactosidase active fractions were pooled and loaded on another 0.2M phosphate buffer (pH 7.4) pre-equilibrated column containing G-75 Sephadex followed by elution with the same buffer to obtain twenty five fractions and assayed for  $\beta$ -Galactosidase activity and total protein.

**Assay of  $\beta$ -Galactosidase and Total Proteins:**  $\beta$ -Galactosidase activity was assayed by monitoring the release (hydrolysis) of methylumbelliferone (MU) from 4-methylumbelliferyl-  $\beta$ -D-galactosides (MU-gal) in the presence of the enzyme. The assay protocol involves incubating 25 $\mu\text{l}$  of 1mM MU-gal and 100 $\mu\text{l}$  of the enzyme for 60 minutes at 37°C and the mixture adjusted to pH 10 with 0.1M NaOH. The absorbance was then read at 360nm.  $\beta$ -Galactosidase activity was defined as the amount of enzyme required to hydrolyse one (1.0)  $\mu\text{mol}$  of MU from MU-gal per minute under the described assay condition. Total proteins were determined by the Biuret method using Bovine serum albumin as standard.

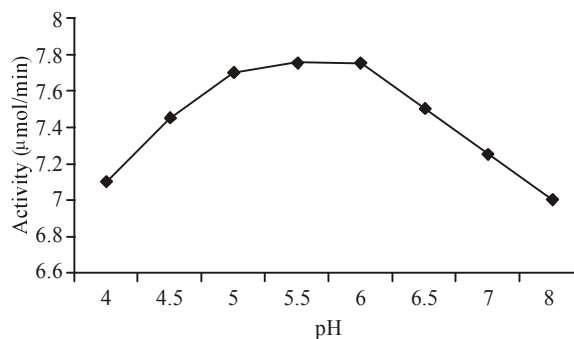
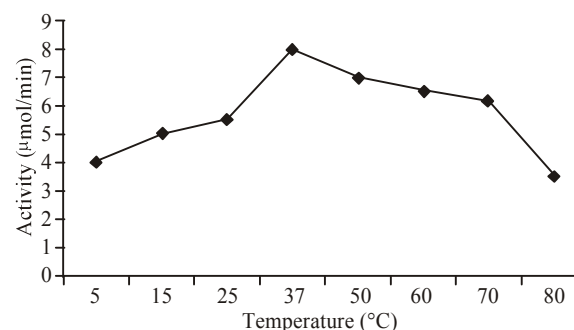
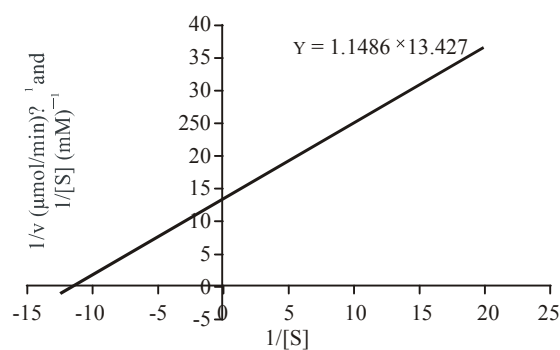
**Temperature and pH Dependent Studies:** A pH dependent assay of the enzyme was performed using 50mM acetate buffer pH 4.0-6.0 and 50mM phosphate buffer pH 6.5-8.0. The temperature-dependent study was conducted by assaying the enzyme activity at varying temperatures between 5 and 80°C.

**Initial Velocity Studies:** Kinetic experiments were performed by assaying the enzyme activity with varying concentrations (0.05-1.0mM) of the MU-gal substrate. Initial velocity values obtained were then used to plot Lineweaver-Burk's plot to determine the  $K_m$  and  $V_{max}$  of the enzyme towards the substrate used.

**Effect of Some Divalent Cations:** The enzyme assay described above was conducted in the presence of the following divalent cations;  $\text{Ca}^{2+}$ ,  $\text{Co}^{2+}$ ,  $\text{Mg}^{2+}$  and  $\text{Fe}^{2+}$  at 10mM final concentrations using their chlorides salts.

## RESULTS

Table 1 summarizes the results of the purification profile of  $\beta$ -Galactosidase from *T. evansi*. No enzyme activity was detected before the addition of the Triton

Fig. 1: Effect of pH on *T. evansi*  $\beta$ -galactosidase activityFig. 2: Effect of temperature on *T. evansi*  $\beta$ -galactosidaseFig. 3: Lineweaver-Burk's plot for *T. evansi*  $\beta$ -galactosidase

X-100. The enzyme was purified 6.9 fold with a 33.33% yield after the gel filtration on Sephadex G-75. The pH activity profile of *T. evansi*  $\beta$ -Galactosidase was a typical bell shaped curve with optimal activity at pH 6.0 (Fig. 1) while the relationship between temperature and enzyme activity was broad with optimal activity at 37°C (Fig. 2).

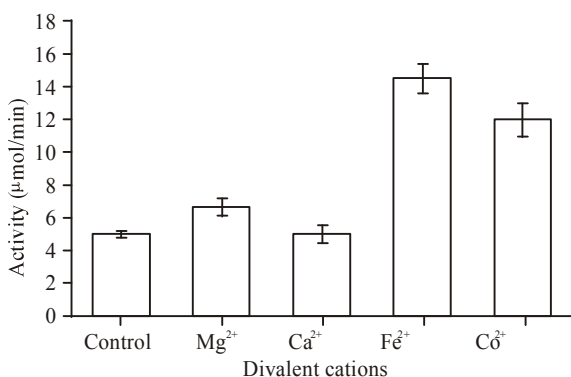


Fig. 4: Effect of divalent cations on  $\beta$ -galactosidase activity

The computed kinetic parameters of the enzyme using Lineweaver- Burk's plot revealed a  $K_m$  of 0.08mM and  $V_{max}$  of 0.07  $\mu\text{mol}/\text{min}$  with MU-gal as substrate (Fig. 3). In the analysis of the effects of some divalent metal cations on the activity of the enzyme,  $\text{Fe}^{2+}$  and  $\text{Co}^{2+}$  were found to activate the enzyme by about 3 folds and a slight activation by  $\text{Mg}^{2+}$  while  $\text{Ca}^{2+}$  caused a mild decrease in the activity of the *T. evansi* enzyme (Fig. 4).

## DISCUSSION

$\beta$ -Galactosidases have been reported in bacteria, viruses and some protozoans [4, 11-13]. So far only *T. cruzi* [8, 9] and *T. congolense* [6, 7] are known to have  $\beta$ -Galactosidase but is yet to be characterized. This report is the first to attempt the purification and characterization of  $\beta$ -Galactosidase from trypanosomes.

The addition of Triton X-100 had effect on *T. evansi*  $\beta$ -Galactosidase activity, indicating that a membrane solubilization is required for the catalytic activity of the enzyme; thus *T. evansi*  $\beta$ -Galactosidase could be a glycosylphosphatidylinositol-linked (GPI) membrane bound enzyme. Sialidase (neuraminidase) from *T. evansi* have been reported by Nok *et al.* [3] to be a GPI linked protein. It will not be surprising that both enzymes because of this common ground exhibit some sort of synergistic role in the pathogenesis of *T. evansi* infection as the sialidase hydrolyse sialic acids from erythrocyte surfaces during the course of the parasites infection exposing  $\beta$ -Galactosyl epitopes on the asialo-erythrocytes surfaces. The increase in specific activity of the crude  $\beta$ -Galactosidase after the two purification steps could be attributed to the removal of other synergistically interacting proteins.

The pH dependent activity profile revealed optimum activity at 6.0. An acidic pH optimum region contributes

to the enhancement of the enzymatic reaction of the *T. evansi*  $\beta$ -Galactosidase by general acid catalytic mechanism. The control of intracellular concentration of  $\text{H}^+$  is a very essential pathological character of *T. evansi*. Indeed, the parasite has a very sensitive pH intracellular pool which is released by ionophore nigericin [14]. *Trichomonas foetus*  $\beta$ -Galactosidase has pH optimum at 5.8 [4] whereas both *Propionibacterium acidipropioni* and *Bifidobacterium longum*  $\beta$ -Galactosidases have been shown to have pH optimum at 7.0 [12, 15]. Thus, the contrasting pH profile could be organism related. The *T. evansi*  $\beta$ -Galactosidase was optimally active at 37°C which is a favorable temperature for parasitic life in the mammalian host. Optimal activities for *P. acidipropioni*, *B. longum* and *Tilapia nilotica* intestine  $\beta$ -Galactosidases [12, 15, 16] have been reported to be within the range 37-50°C. It therefore appears that optimal activity at 37-50°C is a common property of  $\beta$ -Galactosidases.

The fairly low  $K_m$  value of the enzyme is an indication of moderately high affinity of the enzyme for the substrate. The methylumbelliferone component could contribute to the enhanced activity of the enzyme by general acid catalysis and proximity of the participating species. Moreover, the  $V_{max}$  of 0.07  $\mu\text{mol}/\text{min}$  may imply that at the end of 24 hours post *T. evansi* infection, at least 100  $\mu\text{moles}$  of  $\beta$ -Galactosides would have been excised from the host RBC and/or other glycoconjugates. Cleavage of  $\beta$  galactosides which links sialic acids and RBC membrane could lead to the liberation of both sialic acid and the  $\beta$ -Galactosides which consequently accelerate the death of such erythrocytes leading to anemia observed in the course of the disease [3].  $\text{Fe}^{2+}$  and  $\text{Co}^{2+}$  were found to activate the enzyme by about 3 folds whereas  $\text{Mg}^{2+}$  slightly activated the enzyme. The presence of these ions could modulate the activity of the enzyme during the course of *T. evansi* infection.

We have therefore concluded that the bloodstream form of *T. evansi* expresses  $\beta$  galactosidase and was characterized herein. We are presently working on the role (if any) of *T. evansi*  $\beta$ -Galactosidase in causing anemia.

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