



CHARACTERIZATION OF TRANSITION STATE REGULATOR *AbrB* IN BATCH CULTURES OF *Bacillus thuringiensis* USING POLYCLONAL ANTIBODY ANTI-*AbrB*

Astrid Magdalena Lozano Goné¹, Jabel Dinorín Téllez Girón¹, María Eugenia Hidalgo-Lara², Víctor Eric López y López¹.

¹CIBA, Instituto Politécnico Nacional Carretera Estatal Santa Inés Tecuexcomac-Tepetitla Km. 1.5, Tepetitla, Tlaxcala. C. P. 90700. ²CINVESTAV-IPN. Email: vlopezyl@ipn.mx

Key words: *AbrB*, *Bacillus thuringiensis*, transition regulator.

Introduction. *AbrB* is one of the most important regulators of the transition state. It is involved in the regulation of various cellular functions, also has the ability to activate, suppress or prevent the inappropriate expression of various genes⁽¹⁾. From a Bioprocess point of view, *AbrB* might be influence the synthesis of many industrial products synthesized by the *Bacillus* genera. However, most of studies on *AbrB* are made in flask cultures. The characterization *AbrB* in *Bacillus thuringiensis* (*Bt*) has not been reported. The aim of this study was characterized the transition state *AbrB* in batch cultures of *Bt*. We used a recombinant *AbrB* expressed in *E. coli* as antigen for the production of polyclonal antibodies anti-*AbrB* obtained from rabbits.

Methods. *abrB* gene of *Bt kurstaki* HD73 was cloned with vector pJET and *AbrB* protein was expressed in the vector pQE30Xa in *E. coli*. The polyclonal antibody was purified by affinity column with protein A *S. aureus* and tested for reactivity of the immunoglobulin (IgG's) obtained (dilution of 1:200 to 10). Serial dilutions of the recombinant protein *AbrB* (0.01 10 µg/ml) was tested. *Bt* pHT1kAc was grown in clean medium (CL) based on glucose and soy peptone and MHS medium based on glucose and soybean meal. Medium. Batch cultures were performed in 7-l reactor with 4-l of medium culture at 600 rpm, 30°C, 1 vvm. Cellular counts, spores and *cry1Ac* expression were quantified. *AbrB* protein synthesis was followed by ELISA and Western blot using the recombinant *AbrB* as positive control.

Results. The purified *AbrB* and In order to verify the immunoglobulin isotype once purified, was analyzed (Fig. 1).

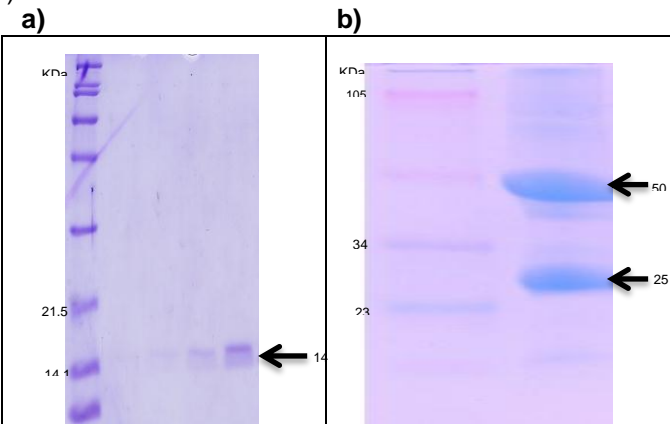


Fig.1 Displays denaturing polyacrylamide electrophoresis (15%), a)Purified protein *AbrB*, b)Purified immunoglobulins.

According to the presence of bands with molecular weights of 25 and 50 kDa, we verified the isotype γ (gamma) of the obtained IgG's.

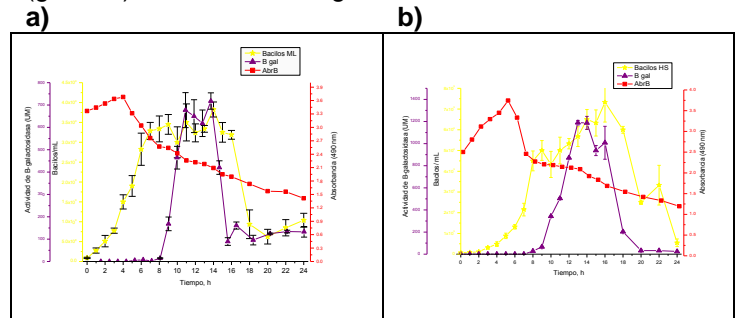


Figure 2. Kinetics of growth, *cry1Ac* expression and *AbrB* synthesis in batch cultures of *Bt* pHT1kAc. *AbrB* synthesis was evaluated by ELISA. The purified antibody was used at a concentration of 25 µg/ml. a) CL, b) MHS.

AbrB synthesis in batch cultures were detected from the beginning of fermentation became maximal at 4h and 5 h for ML and MHS respectively. The changes in expression profiles may be attributable to the medium composition and availability of nutrients. Maximum *AbrB* synthesis coincided with the middle of the exponential growth⁽²⁾. Other reports mentioned that *AbrB* expression became maximal from the late exponential phase until de end of transition state⁽³⁾.

Conclusions. The *AbrB* protein expression occurred from the beginning and middle of exponential growth. Hence, *AbrB* is a regulator of the transition phase from latency to exponential phase, but not a transition regulator from the exponential phase to stationary phase as reported. Now, we are conducted the homologous recombination of *Bt* to evaluate the effect of *AbrB* overexpression on metabolism, *cry* expression and sporulation.

Acknowledgements. CONACYT CB83057, scholarship 250617. Yolanda Medina Laboratorio de Anticuerpos Monoclonales-INDRE.

References

1. - Fisher, M.A. Strauch, M.R. Atkinson, L.V. Wray Jr. 1994. Modulation of *Bacillus subtilis* catabolite repression by transition state regulatory protein *AbrB*, *J.Bacteriol.* (176): 1903-1912.
2. - M. O'Reilly, K.M. Devine 1997. Expression of *AbrB*, a transition state regulator from *Bacillus subtilis*, is growth phase dependent in a manner resembling that of *Fis*, the nucleoid binding protein from *Escherichia coli*, *J. Bacteriol.* (179): 522-529.
3. - Allison V. Banse, Arnaud Chastanet, Lilah Rahn-Lee, Errett C. Hobbs and Richard Losick.(2008). Parallel pathways of repression and antirepression governing the transition to stationary phase in *Bacillus subtilis*. *Proc. Natl. Acad. Sci. USA* 105: 15547-15552.